

SEQUENCE LISTING

<110> Habben, Jeffrey E.
Zinselmeier, Christopher
Tomes, Dwight
Abbitt, Shane
Helentjaris, Timothy G.
Niu, Xiaomu

<120> Modulation of Cytokinin Activity in
Plants

<130> 0803R

<150> US 60/460,718

<151> 2003-04-04

<150> US 09/545,334

<151> 2000-04-07

<150> US 60/129,844

<151> 1999-04-16

<160> 39

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1919

<212> DNA

<213> Agrobacterium tumefaciens

<220>

<221> CDS

<222> (690)...(1411)

<223> ipt

<400> 1

```

ggatcccggtt acaagtattg cacgttttgt aaattgcata ttaatgcaat ctggatgttt 60
aataacgaat gtaatggcgt agaaatatgt attttattgt atttatcttt cactatgttg 120
aagtttgcaa taatatgcta atgtaaaatt aaaaaattat gtactgccgc atttgttcaa 180
atggcgccgt tatttcaaaa atatctttga ttttgttacg aggacaacga ctgcaggaag 240
taaataaaaag acgctgttgt taagaaattg ctatcatatg tgcccagcta tagggccatt 300
taagttcaat tgtgaaatag ccgcccttat tttgacgtct catcaaatca aatattaaaa 360
aatatctcac tctgtcgcca gcaatgatgt aataaccgca gaaaagtgag agtaaatcgc 420
ggaaaaacgt cgccgagtgg catgaatagc ggccctccgta ttgctgattt agtcagcttt 480
atttgactta aggggtgccct cgtttagtgac aaattgcttt caaggagaca gccatgcccc 540
acactttgtt gaaaaacaag ttgccttttg ggaagaacct aaagccactt gctcttcaag 600
gaggaatatc gaggaagaga atataacagc ctctggtaca gacttctctt gtgcaaaaat 660
caatttgtat tcaacatadc gcaagaccg atg gat cta cgt cta att ttc ggt 713
Met Asp Leu Arg Leu Ile Phe Gly

```

1

5

```

cca act tgc aca gga aag aca tcg act gcg ata gct ctt gcc cag cag 761
Pro Thr Cys Thr Gly Lys Thr Ser Thr Ala Ile Ala Leu Ala Gln Gln

```

10	15	20	
act ggc ctc cca gtc ctc tcg ctc gat cgc gtc caa tgc tgt cct caa			809
Thr Gly Leu Pro Val Leu Ser Leu Asp Arg Val Gln Cys Cys Pro Gln			
25	30	35	40
cta tca acc gga agc ggg cga cca aca gtg gaa gaa ctg aaa gga acg			857
Leu Ser Thr Gly Ser Gly Arg Pro Thr Val Glu Glu Leu Lys Gly Thr			
	45	50	55
act cgt ctg tac ctt gat gat cgc cct ttg gta aag ggt atc att aca			905
Thr Arg Leu Tyr Leu Asp Asp Arg Pro Leu Val Lys Gly Ile Ile Thr			
	60	65	70
gcc aag caa gct cat gaa cgg ctc att gcg gag gtg cac aat cac gag			953
Ala Lys Gln Ala His Glu Arg Leu Ile Ala Glu Val His Asn His Glu			
	75	80	85
gcc aaa ggc ggg ctt att ctt gag gga gga tct atc tcg ttg ctc agg			1001
Ala Lys Gly Gly Leu Ile Leu Glu Gly Gly Ser Ile Ser Leu Leu Arg			
	90	95	100
tgc atg gcg caa agt cgt tat tgg aac gcg gat ttt cgt tgg cat att			1049
Cys Met Ala Gln Ser Arg Tyr Trp Asn Ala Asp Phe Arg Trp His Ile			
105	110	115	120
att cgc aac gag tta gca gac gag gag agc ttc atg agc gtg gcc aag			1097
Ile Arg Asn Glu Leu Ala Asp Glu Glu Ser Phe Met Ser Val Ala Lys			
	125	130	135
acc aga gtt aag cag atg tta cgc ccc tct gca ggt ctt tct att atc			1145
Thr Arg Val Lys Gln Met Leu Arg Pro Ser Ala Gly Leu Ser Ile Ile			
	140	145	150
caa gag ttg gtt caa ctt tgg agg gag cct cgg ctg agg ccc ata ctg			1193
Gln Glu Leu Val Gln Leu Trp Arg Glu Pro Arg Leu Arg Pro Ile Leu			
	155	160	165
gaa ggg atc gat gga tat cga tat gcc ctg cta ttt gct acc cag aac			1241
Glu Gly Ile Asp Gly Tyr Arg Tyr Ala Leu Leu Phe Ala Thr Gln Asn			
	170	175	180
cag atc acg ccc gat atg cta ttg cag ctc gac gca gat atg gag aat			1289
Gln Ile Thr Pro Asp Met Leu Leu Gln Leu Asp Ala Asp Met Glu Asn			
185	190	195	200
aaa ttg att cac ggt atc gct cag gag ttt cta atc cat gcg cgt cga			1337
Lys Leu Ile His Gly Ile Ala Gln Glu Phe Leu Ile His Ala Arg Arg			
	205	210	215
cag gaa cag aaa ttc cct ttg gtg ggc gcg aca gct gtc gaa gcg ttt			1385
Gln Glu Gln Lys Phe Pro Leu Val Gly Ala Thr Ala Val Glu Ala Phe			
	220	225	230
gaa gga cca cca ttt cga atg tga ta gattgcacca gttttgtttc			1431
Glu Gly Pro Pro Phe Arg Met *			
235			

```

agacttgctg ctatttgaat aagatgttcg ttctttgttg tgttggtgtg ttgtgataga 1491
ggcaagtggg ttgaaacttg tttttactgg tttattttca gtctcttgga cgatgtttta 1551
caaataataat attgtgaaaa ttgtgggttt atattcgtag aacgaaataa atggtaagta 1611
tagccgttat caaaatttag caaaaattgt taaaggttct tttatgcggt gaggttgctg 1671
acttttcatc attgtcgcgt aaggagttac ggatatccat aactgtaaaa acgccgcaga 1731
atttacgggt ggtgcattta gtttgccgtt caacatgatt ttggcaatag ttggtaacca 1791
agcactagcc aaccgttcga taatcactta atcgatggaa ccgttcagct ttccttcgtg 1851
aggctgctct tgatgatgag ctgccgtcta gtttttataa cgccgggtta cgcattatag 1911
acaagctt                                     1919

```

<210> 2

<211> 239

<212> PRT

<213> *Agrobacterium tumefaciens*

<400> 2

```

Met Asp Leu Arg Leu Ile Phe Gly Pro Thr Cys Thr Gly Lys Thr Ser
 1          5          10          15
Thr Ala Ile Ala Leu Ala Gln Gln Thr Gly Leu Pro Val Leu Ser Leu
          20          25          30
Asp Arg Val Gln Cys Cys Pro Gln Leu Ser Thr Gly Ser Gly Arg Pro
          35          40          45
Thr Val Glu Glu Leu Lys Gly Thr Thr Arg Leu Tyr Leu Asp Asp Arg
          50          55          60
Pro Leu Val Lys Gly Ile Ile Thr Ala Lys Gln Ala His Glu Arg Leu
65          70          75          80
Ile Ala Glu Val His Asn His Glu Ala Lys Gly Gly Leu Ile Leu Glu
          85          90          95
Gly Gly Ser Ile Ser Leu Leu Arg Cys Met Ala Gln Ser Arg Tyr Trp
          100          105          110
Asn Ala Asp Phe Arg Trp His Ile Ile Arg Asn Glu Leu Ala Asp Glu
          115          120          125
Glu Ser Phe Met Ser Val Ala Lys Thr Arg Val Lys Gln Met Leu Arg
          130          135          140
Pro Ser Ala Gly Leu Ser Ile Ile Gln Glu Leu Val Gln Leu Trp Arg
145          150          155          160
Glu Pro Arg Leu Arg Pro Ile Leu Glu Gly Ile Asp Gly Tyr Arg Tyr
          165          170          175
Ala Leu Leu Phe Ala Thr Gln Asn Gln Ile Thr Pro Asp Met Leu Leu
          180          185          190
Gln Leu Asp Ala Asp Met Glu Asn Lys Leu Ile His Gly Ile Ala Gln
          195          200          205
Glu Phe Leu Ile His Ala Arg Arg Gln Glu Gln Lys Phe Pro Leu Val
          210          215          220
Gly Ala Thr Ala Val Glu Ala Phe Glu Gly Pro Pro Phe Arg Met
225          230          235

```

<210> 3

<211> 2085

<212> DNA

<213> *Zea mays*

<220>

<221> promoter

<222> (1) ... (2085)

<223> zag2.1

<400> 3

```
agcttcgtgt gttccttcga tcggtcacag tttgattcct gctcaccaca tattttttgcc 60
gcgtgggagg gaggccacga ctggtggcag aacagcgaga ggcagactac ccttacagcc 120
ttaataactc ttatatcttc tactataaca tcaaaataag acgtagtgtg gtggatatgt 180
tgtctcta at ttagcagcag gtcttgagtt tgattcaca ttcttgcaga tttatttttt 240
gagccataac agggatgagg gcaaaatagg aaatgaacga catgttacct ttaccgcctt 300
aataagtagt agagatatcc agttttatac taattattat tatataaaat gcactgcaca 360
tatattacta ttaccagttt tcttggacat gcacagcaga aaacacgcac acgcagagag 420
gaaaaggaga ggccataaac caaaaggctt taagaatata tgtaaagata tgtctaaatg 480
gctatatctg gttaagcaag ataacagggc tctggtcatc agtagtagtg gcctttttgcc 540
cttgccctc atctctctca cactctctt ttctcagcct tgcttccgat cgatggatcc 600
catcccactg ccatagtgcc atcctttctt tcccttgcgc gcattgccta gccggcggc 660
cggcctgcta ttaaaccact ttacccccct tctcggtcac gctcgacgca gctccctttt 720
ccttgcttgc ttattgcaag tctctgcaag aacctgctag agaggaacaa ggtagaatag 780
tatcgctttt tccatctaga ggttatctct ttttacatga aaaatttcag ccgtattttc 840
gttctccata tatcagtcct gcgataatat aaatagcgcg gtcttgtgtg atccggcata 900
tgtatagttc ctactaactg atcgagatcg ctctcgtttg tactttctcc ctttgaggaa 960
agagttcccc tttttctgtg cttcaaattc ttgtaaggaa aacctgcct gcctgccagc 1020
ttcttctgct acttggtatg tgattcttat ttgcttactt gatttccgtt ttttttctt 1080
gctttctata tgtatgtatc tgggctgtct tccctgcgt ctcgttacta cgtactaagc 1140
tttggaaggt ttcaactctt tgtatacgat gaggtttctg cccctagtag cagatccgcg 1200
cacgactaga tgtttgagga aaagaaaagg gcaagacgct atatatatat gcagcagcga 1260
gtcgcacata tatccagttt tccaatctgc ctcttgcttt atgataattc aacttgcgct 1320
gattatatcc ttggctacct agctagaaat gtctaattaa actttgtttg ctagctagat 1380
tttggtgctt cttttcgcat ctgatctttt tatctcttct gagtgctccg caaagccttc 1440
cagtgttgaa gaagctgctg gaagaagaga tgagctttct cttgaaggaa aaagagatga 1500
tcattgccgg tttgttggtt tttcgtgtt ttttagcttc ttgtcccca tttatattcg 1560
cgcctaatac acgagcccg agatcttggt ttcttgtggc tggttttgtt ggatctcgat 1620
ctcggttacg ttacatgag tcttgctgcc taacatacat ctgtgttctt tttctaggct 1680
gcgagaaact taactgatcg agtctgtctg gcaggcatcg atctatccag tcgtcagttc 1740
gtcacatccg ctttttcgta tatatcatct tcagattttg tccatctgtc aaatcatgga 1800
aaatctgtcg tctttgcttg tattctcttc tgttattcct gctgctccg gccggaccaat 1860
tcttgaatcg acccggtgtc ctattccctt ttgttagaca gccc aaatcg cttgctcgat 1920
cgtagtgtac tgtactactg cggctagcta gatcttccaa gctagctata gttcgccggg 1980
ccctttgatc tgcttcacag aacatatata acacttgaac tcttttacgc ttatgagaaa 2040
acttgctgct tgctgcttcc agctgggtatc gtcgccagcg gatcc 2085
```

<210> 4

<211> 344

<212> DNA

<213> Cauliflower mosaic virus

<220>

<221> enhancer

<222> (1) ... (344)

<223> CaMV35s

<400> 4

```
tctagaaatc cgtcaacatg gtggagcacg acactctcgt ctactccaag aatatcaaag 60
atacagtctc agaagaccaa agggctattg agacttttca acaaagggtg atatcgggaa 120
acctcctcgg attccattgc ccagctatct gtcacttcat caaaaggaca gtagaaaagg 180
aaggtggcac ctacaaatgc catcattgcg ataaaggaaa ggctatcggt caagatgcct 240
ctgccgacag tgggtcccaa gatggacccc caccacgag gagcatcggt gaaaaagaag 300
acgttccaac cacgtcttca aagcaagtgg attgatgtga tgct 344
```

<210> 5

<211> 2198

<212> DNA

<213> Zea mays

<220>

<221> promoter

<222> (1) ... (2198)

<223> ZmMADS

<400> 5

```
ccttttttctt tttctccaca acatgaacct tactagaaca ctgccccact taaaagaatg 60
agggtagaac tcttgaatct tagggatttg aactccttgc agtacctcat aacaaggggtg 120
ttacatgtcc ttcttctgct gttgctgctt gagcaggata tagagagatg accgacaccg 180
ggttgatctt gggacaacct tcttctcctc ttttcttcgt tgttttcttt tctatttctca 240
ctaccttttt ctttctcttt gttcttccca ctggaggatt ctatcaaaaa gtattaccat 300
catacagagg aggaacccga agactatgaa ccatgtacaa cagtcttcaa cccaagaatc 360
accaagcatt gtgatcttag gggcgaggga gtggaaaatg gagttgcttg tgatttggca 420
gaggggaattt tatcaggagt gttttgcttt gagtggaatg ggaactgagg gagttggttg 480
ggggggggggg tttataggcg agtgggagtg ctcggtgctg gagtggtggtg atggaacagg 540
tgacatgagg tagcaggtcg atggaggggg gctgttgccg gcgatgatgg cggcggtggg 600
tgcgctgcaa aggagggcgt ggggcggtgg tagtgcgcat ggaggcgggc acgcgtgctg 660
ggggcacaag tgagtgggtg ggtcgatgac cctgatgttt gtggtctctg gttccaagaa 720
tctttgtctc tctttatgat aataacttct tttgtcgtcc ttttctgttt actttgactc 780
aggggcagtg ctttgattct cacggtcggt ctttttgact gagtgactgg acatgtttct 840
tctgtagcat tgtacaacat gtactttgtg caagctacaa ggccacattt tttgaagcat 900
agattctttc ccccaaacaa tttatacaaa tatgcaaggc tacacttctt gtatttctat 960
aacattgtac attcatgaca gaggtcctaa agcttgtaaa ttttgtgcag gtttaattca 1020
tgtaaagttc ccttgtagag tcatgacaac atcgtactat aaaattattc taaaaaaccc 1080
acacatgacc cccatgttat ttggtgacaa tacagaaacc acacatctag tgatgatata 1140
acactgtaca gaagccacaa attataatat ataaaacact atacaaagta tccaaataaa 1200
gcctaataag tatggagggt aacctgaatc tttcctaata ataatagaata atctacaata 1260
atgatttggt tggaacaaaga gaattaaacg gtattgagtg ggctaaaatt ccttggttatt 1320
caaaaccctc aatcacagtt tctccgaggg aaaaagaaac aggggaggag actcaggctg 1380
ttcacaatag ggatttcata tcgctctttc caacaatgcc acatcatcaa aagtgttatg 1440
aaactaaaaa tgaaataata cttctcaatg caaactttca ttttcataga ttaatatact 1500
aattaaatga tgcaactaaa taaccaatag atgttagtaa aatatggtaa gattaaacaa 1560
accactatca atggacattt cacatagttt ccaagacttt gaaaacgggt tgacatgatt 1620
tcatccacat caaactaatt ttatctctga aaccatttca ttttaaataa tatggcataa 1680
cgtccaaaat gctgacgtga cataccatta aatgtgcatg aaactcccat aaaactttta 1740
ttgataatag cctcacagac atccggtcct acaccgtgtt ggacccatca gccagacgcc 1800
ctgcagcaaa cgcgacgttt gacttgccat ctgctccct tgtgcccga cgcacctgga 1860
aggctggact ggaactggaa caagcaaaat ggaaaaaac atatctcacc actgaaccgc 1920
acccttcgag ccacgcccag gctcgaccaa tccctgcccc gcgcgccttg acgagcgcct 1980
cactcgaacg ccggcctcgc taggcccctc cttctggccc gcaataacga tccccgtcat 2040
gatccgacgg tctagctgcc tccacgccgc tccaaaaccc ccgcgtccaa tcaaaacacg 2100
acagcggggc gagcgaaccc accgtggttt cgccaaaccg ctttcccttc catctaaaac 2160
cgccccctcc cttcctcttc tcctagctct cttgcctg 2198
```

<210> 6

<211> 1470

<212> DNA

<213> Zea mays

<220>

<221> promoter

<222> (1) ... (1470)

<223> ckl1-2

<400> 6

```
gagctcgccc ttgcatgctt gagtcatatc ttggaaaaaa aaactgtaac ttaaagtatg 60
```

atctatatat	ggattatattg	gatgggatgt	catttttcgta	tcaccaacca	aaattacagt	120
ttggtcgtgc	gtagaaattc	tacctactag	ctgaaacaac	ggctgctatg	tataactact	180
ggtactggaa	agaatattag	tcattgactc	aaaattagaa	tgcattgtgta	agtcattgcgt	240
gctaattttgt	tctatcagca	ttcggcgaat	tccgaagtcc	gtacgtggtg	ttcgtggagg	300
agaggaaaac	atcagaaatg	acaaaactag	acggcgtgtg	cttctacact	gaattcatca	360
acattttgttt	tactttttact	agagaatggc	atcagatgga	aaaccgctga	aaaaacaaga	420
aaacaattgg	accccaaata	tgtacagacg	ctagctatag	ccagccacac	tgaagttgac	480
atgcggaac	tagctaacca	ccttctctga	aacactaaca	tttgtacctt	ggctcgtgtaa	540
gtgtagttag	taacgtatgt	tgacgcgact	taccgaacaa	aaatataatt	gtcccaatca	600
agctagggac	gattgtttgt	ttccaaaatg	ttgccatttg	cttaatcaat	cctatatattga	660
ttcatggctg	ttaaggtgag	ataaagcgac	aagaaatctc	tctctatata	tatatataag	720
atcccgaagg	ctagegcacat	ttttgatagc	aaaatatgag	aagttggcag	gttctggtag	780
caaatcaaat	aatatggcca	gaataatcgt	ggctagcttg	attaaacctt	cagcttggtg	840
tatttttgaa	gtcgaccaac	cagctgggcc	ggggctcgtc	gtagtaccaa	aattacagcc	900
tgcttccttc	gtcgtcctgt	acgtaatgca	gtacagctgt	ctgtctagta	gagacgattt	960
tgagcaggca	cacacattaa	gtgataacat	aaaagacggc	ttcattttat	ttcataacca	1020
aacgatattg	tcaacacaca	cctatagcta	ccaaatttgt	acaactattt	agtgcgaaaa	1080
ctatttcatt	ctcaagaatt	gatcgcttat	atttattatt	acagggtttt	aatgtataa	1140
atagcgtata	ttgcatggca	aaagggggta	ataattaggc	aggactatat	atataatagt	1200
tttttttct	ttaaattcct	gggaggatgg	taaagttggt	aactaggcac	cttggtgcgca	1260
tatttttctg	tggtcaaaca	gaataaaact	agacgggatg	cagaattttt	ttttccttgg	1320
aaagcagctc	atctctgtgt	tcgagtacgt	aattgaagaa	gtatgtgatc	gcactacacc	1380
tacacgtatg	tgccgccgta	tccgtcctat	atatatacgg	ggtgcaatca	cctagttacc	1440
aaacactcac	acataagggc	ggatccatgg				1470

<210> 7
 <211> 960
 <212> DNA
 <213> Zea mays

<220>
 <221> promoter
 <222> (1) ... (960)
 <223> eep1

<400> 7	
tcaaaccggt	catcgtttgt
taagacagct	gtgaggggtg
attagttttt	ttttcttggc
ttcagttatt	cgtgtgtcct
acttgctatg	gatttaatta
aacaagctct	aagggtgata
tttttctaca	gatgggtcag
ttttttaacg	acagggttat
agtgtctact	aattactaac
aagaatctcg	gaacatttaa
cctagctgga	tactacttat
tccaaggtca	aactgaatgc
acgcctgatg	tgatgtaatg
taccagggct	tgcatggaaa
atacacatac	atgcacacca
aaaaatacag	agagagagag
atcatccact	gcttgacttg
gacccaacta	accataaat
aatcattcaa	gttggcatga
agtgattgaa	aaaaattatg
cctccaatcc	atatggattg
tatatgctgc	atgagcagtt
tggtgaatgc	atcagggtcat
acacaccata	tattccctaa
cgtatgctca	ttgacgtctc
ttgatactac	gtgctagctg
aattttctttt	tttgttttct
ttgttacaac	ggtcctctcc
aagagcctta	ttattgtata
tctcattata	ttataaatat
ggactgtcat	ctctcagttg
accctgcaaa	gatatcgacc
60	
120	
180	
240	
300	
360	
420	
480	
540	
600	
660	
720	
780	
840	
900	
960	

<210> 8
 <211> 1224
 <212> DNA
 <213> Zea mays

<220>
 <221> promoter
 <222> (1) ... (1224)
 <223> end2

<400> 8
 tactataggg cacgcgtggt cgacggcccg ggctggtaaa aagtaattga acccaaaata 60
 tcatgggatg tttgggtgaag acagtgatca gtgatttttt tatatctata tatatatcaa 120
 agatacttga ttttctagaa gggtcttttt gttgttttcc cttatgtttt tacgcatgat 180
 gcaattcttt ttgagagggt tccgatgcat tgatgttatt gtattatctc ctatatatag 240
 gtcgacgtac attatgtatt gcaataacca gttaactgga tccagcttcg cttagttttt 300
 agtttttggc agaaaaaatg atcaatgttt cacaaccaa atattttttat aactttttgat 360
 gaaagaagat caccacgggc atatctaggg gtggtaacaa attgcatctt aaatgtttct 420
 tcataaaaaa taaggcttct taataaattt tagttcaaaa taaatacgaa taaagtctga 480
 ttctaactcg attcgatcct taaattttat aatgcaaaat ttagagctca ttaccacctc 540
 tagtcatatg tctagtctga ggtatatcca aaaagccctt tctctaaatt ccacacccaa 600
 ctcatagttt tgcaataaaa tactccgact ccaaaatgta ggtgaagtgc aactttctcc 660
 attttatatc aacatttgtt attttttgtt taacatttca cactcaaac taattaataa 720
 aatacgtggt tgttgaacgt gcgcacatgt ctcccttaca ttatgttttt ttatttatgt 780
 attattgttg ttttctcccg aacaacttgt caacatatca tcattggtct ttaatattta 840
 tgaatatgga agcctagtta ttacacttg gctacacact agttgtagtt ttgccacttg 900
 tctaacatgc aactctagta gttttgccac ttgacctgga cgcgactcta gtattgacac 960
 ttgtatagca aataatgcc aacgacacc tggccttaca tgaaacatta tttttgacac 1020
 ttgtatacca tgcaacatta ccattgacat ttgtccatac acatttatatc aaatatattg 1080
 agcgcatgtc acaaaactga tacaagctg gatgacctc cctcaccaca tctataaaaa 1140
 cccgagcgct actgtaaact actcacaaca caacacatat ctttttagtaa cttttcaata 1200
 ggcgtccccc aagaactagt aaac 1224

<210> 9
 <211> 1433
 <212> DNA
 <213> Zea mays

<220>
 <221> promoter
 <222> (1) ... (1433)
 <223> lec1

<400> 9
 tcctaattctt caaataacca tctcaaaagt tttttaaaac atcttttgag gatatgtatc 60
 ccatagccct agagcgctaa attgactact tttagtcgat taaaaggat tagacatcct 120
 tacaagtcct aagtatcaaa tcaccttcta tcggctatac acaactaacg gaagttatct 180
 ctagtccacac taacttatgt cggtttccgc atggcagatc aaaattagct aacttttgtt 240
 ggctaataag agcaattcca aaagaacgtg taaactaatc tcaaaacaga tattagttaa 300
 gaatagtaat ttttcttact ccaacagttc cctcagtcct ccccaaaaaa ttaagcgttc 360
 cgcacccaca gcctcctctc ggtcgtatct tgggtgtgtt catccctccc caatccattt 420
 ctcaacgtat cagatcatcc accgcctacg acgactgtac agtttgctgc acatatcaca 480
 tttaaaggaa ctgttgaggt acccatcata attcactcct aaaaaatttt agcctgctct 540
 caataatcaa ttgggggggt aaaattttta acatcctttc ggatctaata caacttatgg 600
 aagtttagcta gctctggtcg cgctaacttc tgtcgatcgc ctattagcta atactccatc 660
 tgtcccatta tataaggtat aaccaactct gattcaaaga ccaaaaatat acttaattgt 720
 gtctatacca cttcatcgat gtacgtatgc atagaagag cacatcttat attgtggaac 780
 aagaacaaaa atattggttac gccttatatt ataagacgta gaaatcaatg gtttacaata 840
 gccagaata gatgttttta tttatttctt atatagatgt ttttatttat ttcctatatg 900
 tttcacaata gccttatatt gtgccgaaaa tttaggcaca cgtgccacga acgtctgaaa 960
 tgtactccgc gcgtattacc atgcactacg acgtacgtag gagtatgtac gttgaaccaa 1020
 gcacacatat atctctgaca cagtacaatg atatactaca acaacaacag tactgcccac 1080

ttcatccatt	ttcacgttcc	atcttccgcg	tgtgacaact	cgatcggcca	cgcacgcaga	1140
cgacgacgga	gcagtacttc	acagaatcct	ccgccactcg	tcacaccaac	aggcgcgcg	1200
tggtgcgcat	gcatcatgtg	catgccatcg	tccgtccctt	ggcgtgcctc	ggtagacggg	1260
agctagagta	gtagcctgtg	cttgctaccc	ctgggtcaaca	catcgtagcc	tcctatatatt	1320
aacgtatcct	cacacatcac	aagaacgaca	cacagaaacc	agtagccact	actccatcca	1380
ccacgagcga	gcgagcgata	accctagcta	gcttcaggat	ccagcgagag	ccc	1433

<210> 10
 <211> 820
 <212> DNA
 <213> Zea mays

<220>
 <221> promoter
 <222> (1) ... (820)
 <223> F3.7 promoter

<400> 10	
gagctcaagc	cgcaacaaca aatttcggtg ctcccaagct tcataaaggc tatcttcggc 60
gtcgttgga	tccatggtgg cacagaatcg agttgatgtt gtagctggcg gctaggggtt 120
gaagtggaga	agaggtccgg ctggtggcat cctatcgtct attgaggggt gggtcgggtg 180
gcatcatact	tgatgacaat tgaaagtaat tttaatcaac ttgtcatgag tagtgagtct 240
tttataaaaa	ataagctgaa ataagcacc tttgatgagc ttataggatt atcataatct 300
caaagtctaa	attatataat ttatttagat aagttgcttg ttgtttccc cactagctta 360
tttacattgg	attatataat ctacataaat tataatctca aacaaaaagt ccttaatcag 420
agatcagcga	ggtctcacga gtgagaaggc gagagcttgt ccaaacgagc attttcgggc 480
gtgtgaacac	ccatttcagc aaagccgctg ttgtccagtt cagcgaagcg cattctgcgg 540
ctttggcggtg	accatttctg ctagctcagc actgagaata cgcgtccgct gcagcgttgg 600
cgtacaggcc	ggactacatt agccaacgcg tatcggcagt ggcaaacctc ttcgcttcta 660
actccgctgg	gccaccagct ttgaccgccc cctcccttcc cctccgctac tgctcctccc 720
caccccactc	ccccgcagga gcggcgggcg cggcgggcag gtcgtacccc acatcggcga 780
gcggcgggcg	caccgcccga ggcaaaggca agtctagaac 820

<210> 11
 <211> 26
 <212> DNA
 <213> Zea mays

<400> 11	
gtcagtggtg	taaaagcact tctggt 26

<210> 12
 <211> 26
 <212> DNA
 <213> Zea mays

<400> 12	
tgcgccagaa	gaagcagcag gaagat 26

<210> 13
 <211> 26
 <212> DNA
 <213> Zea mays

<400> 13	
aagcttaggg	tacctcaaac cggcca 26

<210> 14

<211> 34
 <212> DNA
 <213> Zea mays

<400> 14
 ccatgggtcga tatcttttgca gggtagggat ctct 34

<210> 15
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Clontech AP1 primer

<400> 15
 gtaatacgac tcactatagg gc 22

<210> 16
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Clontech AP2 primer

<400> 16
 actatagggc acgctgtgt 19

<210> 17
 <211> 1679
 <212> DNA
 <213> Zea mays

<220>
 <221> promoter
 <222> (1)...(1679)
 <223> tb1 promoter

<400> 17
 gcggccgcct acctaataga tatgtatcac tctctctca cttcggctat aaaagagagg 60
 gatagagaaa catagaatgg gtttcgaaaa aaactctttg actctctaaa tagaaacaag 120
 aggaaaggga agttagttgg tcattatctt tgttgatgcc tcaacatgta attttcttcg 180
 ccattgtatt tctcaatcca ctatatacaa agagggtata gggatatatat tacacatctt 240
 acggtccgaa cctatattta aattacccat gtattgatgc ctaggcggta tccagcaaca 300
 gagtgtctct agcacgcatc tcttactcta tttatcaact ctccccgaa tacatgttgt 360
 tccttattgt cactggcgga tctacagggt gtcaccctgt agtccgggtac cggcataaca 420
 tattagcttt gtctatttca tgacttcaaa catggttgcaa caacctacag atgcgttcag 480
 tctatctata tacaagagga agaatacaag tgacaaatct aatttgtgaa tataagaatt 540
 attatgctgg ttacataga ataccaaatt atagcacaca tttatcattc cttattgaat 600
 ttctaaaatg atttactga attattcatg catttttaat ttggcatacc ttatagtaaa 660
 attctataac cgctactgct tattgtcatt atgcgacttg gaagacattt tctacctact 720
 gaaagcggtc tgttttttgt gttgtcgaga gtgtgatggg taaccatagt taataatgca 780
 ctggatctat cactactcat acagggtcca tatgcctaata aatggttgga agaccaactc 840
 atctgaccac atctgtccct accatgcttg tacaccacac tacatacatc actcatcact 900
 ggtccttcgt ttcggtaccc tcctcccaaca atgttcaatg tatatactaa tagttctcaa 960
 ataaattcct gtggatgtta caaaaacca cgggtctttg tttcctgaag aagtatttca 1020
 tggaggcgcg cacgtccatc gtactgcgtc ctgcagctat ggccgcccc atctggccaa 1080

taaagtact	aggtcacttg	tagccaatag	cgtttcaaca	tgcacacagc	ttttcccca	1140
atagtgcagg	tccttgatt	ctcctccctc	tcctcacct	caaattctcat	ccacacgaac	1200
aggcggcacg	gcagtattcc	tccacagccc	tcctctctat	aagatggcac	agccctctca	1260
ggtaggggcg	agtgtctcac	tctcacatag	taaaaaaaaa	aaaaacgccc	ccaaggttct	1320
taagcacaat	tctctagcta	tcttggtctc	ctacacagcc	tatgcacatg	agcccatgcc	1380
tctcctctcc	ttgcgcctgc	atagagaggt	ggtatgatca	cctggaaagt	ttttaactct	1440
ctctctctct	ctctctctct	ctctctctta	caagcctaga	ccttatgcat	ggtcggacgg	1500
acacatctga	tcataggaca	tatgagtagg	ccacactcct	cctgcccctc	tctcgtagag	1560
atcaacacac	actgctctta	gtgccaggac	ctagagaggg	gagcgtggag	agggcatcag	1620
ggggccttgg	agtcccatca	gtaaagcaca	tgtttccttt	ctgtgattcc	tcaagcccc	1679

<210> 18

<211> 1027

<212> DNA

<213> Zea mays

<220>

<221> promoter

<222> (1)...(1027)

<223> eep2 promoter

<400> 18

gtaaagttac	aatttatatat	caaattgctag	ctactagtcg	ggagaaaacc	aactaagggg	60
atgtttgttt	gggattgtaa	tctgtccaga	atatataatc	caacaaattt	tgaactaaca	120
ctcggttcaa	aattttattag	atttatataat	ccatacatat	tacaatccca	aacaaacacc	180
cctaattcta	aatggtgaga	gtaaaaagcg	ctgtctaata	actttttatca	gctaattttgt	240
ttatcttgag	ctgttaatta	aaccattagt	gaagtttttt	tgggggggtgg	tcgaatagag	300
ctaattctaac	tattagctca	taggatcaag	gccattgggt	taatttcacc	ccactatgac	360
tatgtcccag	taactaaata	ctatatattgt	caccataaac	tttgggaagaa	attagttgct	420
actagaaaga	agatccaaac	ctggaaaaaa	ttagttttcta	ctagaaagca	gatcatgtct	480
gctaccaga	cattgattta	tactccagca	tcaaccaacc	ccgtacttgt	tactacaaaa	540
ttggaagaaa	ttagttgcta	ctagaaagta	gataattttct	gccaccagat	attgattttat	600
aacctagtat	caatctctac	tagccttgct	tccgtcattt	gttgctagat	ataaatgggt	660
ttctttcaca	tatgtgagt	tatatatatg	aaccttgcag	caaccattat	attcggtagt	720
caacaaaagc	cctacagaca	tcgatctctg	atctgagaaa	aaaaatcctt	atatggcgag	780
aattacaatg	gaagcaagca	aggctgtcct	gctcttgatg	gtgacctag	gaagtttgat	840
gattcccgca	tactgtaagt	gcacatcg	caaccatg	catttgaatc	aagttacata	900
ttatacagtt	tcttactagt	agtaaatata	aattgttcgc	ataatgtcaa	caaccttaac	960
ttactgtaaa	aacagtaact	gaatgccctt	attgcatgca	gctcggaacc	ttgttcgttt	1020
tctgccc						1027

<210> 19

<211> 723

<212> DNA

<213> Zea mays

<220>

<221> promoter

<222> (1)...(723)

<223> trxl or thxH promoter (thioredoxin H)

<400> 19

gcccttacta	tagggcacgc	gtgggtcgagc	gcccgggctg	gtactctctg	gtactgagtt	60
agatttggtg	aatgttaata	catatatact	tttaataaaa	ttacttttta	agacaaaatt	120
gatgcactgg	cgttcattgg	cggctgtggt	aacaaaaccg	aagtggaagt	agcccgttcc	180
actggaggtt	ggcttaagt	cacatgcagt	gaaaataacg	ttccacttgc	gattcattta	240
acacaactgt	cagtataaat	agtttttttt	attggcggtt	gatttaggtg	aaccccaagc	300
gaaaatatat	ttacacatgc	ggttttttta	gccgtgctca	cctattttatt	ttcagtgtgc	360

ttaactgaaa	ctgtcgggat	aaatTTTTTgc	gtgccatcag	tttagagcac	ttatctactg	420
actTTTTTTTT	tcaagtatcg	tacggatttt	gcaccacgtc	gacgaccgtc	gataacgagg	480
cacgccgatc	tagagagctc	gaagacctgg	gaatggcaca	ggggaccggc	cggagcccg	540
cggcgccatg	caagctgcct	cgatcgcggg	cctcgacctc	agtagcccg	ccctgtcgcg	600
cgccagtcgc	tcgctgcgcc	tataaaagcc	gcccgcggct	cgcgtaggct	accagcgcaa	660
aactctgcca	agggttcgg	atccacacc	gaggaaagga	gaagagagg	tcggaatacc	720
atg						723

<210> 20
 <211> 1626
 <212> DNA
 <213> Zea mays

<220>
 <221> promoter
 <222> (1) ... (1626)
 <223> Zm40 or Mze40-2 promoter

aagcttagct	agatcatttg	taagaatgca	acttgttcat	atagcatggc	tacagcctac	60
atcatctgaa	atagacctgt	ttataggata	cctaagctca	attcacccta	tatctaaaac	120
ctacgaggcc	taaacacacc	cgtcctcaag	aaaacgacca	aaccaaacca	aaccatgcgt	180
ccgtgtcatg	gttttgtaga	cacgtttacg	tatcaattat	agtgttctga	ttttttatat	240
tctcctaatt	atthagagct	aaatTTTattt	ttatgatagc	agagatctaa	atatttttgt	300
tttgattttt	tatatactaa	aatcatctct	acaatattag	agatttttaa	tgctcagaag	360
aatTTTactt	gaattaaaac	ctttactgat	ttttaactaa	aacggagacc	aaaagaaatc	420
tatccaaggc	tgctctaaag	agccttcgtg	tctcgTTTTc	ttatttcaga	cttcactcat	480
cttcttattt	caggctccac	tatataaggt	ggctcttagt	atctttccta	tcacatatcc	540
tattttaaac	tttagtatat	aaaacattat	aattcataat	ataaatcgat	tattttacac	600
gatctcagcc	taaaagcgg	aatatgcacg	ctctgagcat	ggcccaagct	ccacgttaac	660
cgttctgtca	aaaaaaaaa	catctagtct	agaatggaaa	acacacgatt	ttagaagtta	720
ggactagttt	ggcaactcaa	ttttccaaat	gattctcatt	cttttaagag	gatttaattt	780
atTTTTtggt	aaaataggaa	tcactagaaa	ctctattttt	tcaagagaaa	gtaagctatt	840
tttttagaaa	aataaaaaat	cccttaaaaa	atattgttcg	taaattagcc	ctaagatgga	900
ctaaaaatct	ggttttatag	aataggagg	gatcgagcaa	ccgccaatc	tacgcgcaa	960
aaagggtacct	tttcggtgaa	taaacacgac	tgcggcgac	acgatctgat	cgaactcgta	1020
gaataaaaatg	gagcagcgga	atagtgtggg	aggcacaagc	acaggaggag	ctgaaaccga	1080
accgaagtgg	cgaacacgat	ccccactccg	gccggcaccc	gagtgtgcca	gacgtgtggg	1140
gctgatctga	cgagcctgga	agaagaagaa	gaaaaaaaaa	tcctcacgct	cctgcttggc	1200
tccatcgaca	gctcactagc	tgctaccgga	tgctcgcgct	tctgatgcct	ctcgattcat	1260
catccatcgt	tggtggcgcc	ggcgggcgcc	caaaggttct	gattccgcag	cagccaagt	1320
ctcctcctgc	agacgaaaat	gacggcagag	gttggcggtg	atccaggaga	ctcatcagtt	1380
tagtttaata	atgaatctgt	agcaggcgct	tcagtctctc	atcggatgag	cgagcagctt	1440
agcagagcag	gtgggtggtcc	ctggctcgcc	cccgtccatt	ctttcccgcc	cgtcctgccg	1500
tccactccgc	cgctatttta	taccctcct	cgcccaccct	gccatcctca	ccatcgcaat	1560
tcacaagcaa	agcaatcaga	gccaagcacc	caccgtcctc	ctttctttcc	ttcgactcat	1620
caaagc						1626

<210> 21
 <211> 27
 <212> DNA
 <213> Zea mays

<400> 21
 aaacaccttc ggatattgct ccctttt

27

<210> 22
 <211> 27

<212> DNA
<213> Zea mays

<400> 22
tctcgcattt gcagaaacga acaacgt

27

<210> 23
<211> 525
<212> DNA
<213> Zea mays

<220>
<221> promoter
<222> (1) ... (525)
<223> mLIP15

<400> 23
ctttcagcta agtccctgct ccctctcttt ttcttacatt caggtcctcg cagctcctct 60
cttttttctt gtttctttct ttcgatctgc gagccgtcca ggtccagtag tctcctttcc 120
gtgaaggaac tcttgagacc ggccctctcg gtttctctga attcttggtc cccggtccct 180
cctcctgtcc ccgcgtagat ccgtccgtcc gaggagcaca ccgtcccccac ccccatgttt 240
acccaccagt tctctgacg gccgcctgct tccgatgaag ctgagcgtgc tccgtatccg 300
ccgctccac tcttctctcg tcgccttctt ctactgggtc tacgtcttct catgaacgca 360
tcgcccctct ccacctgctg atccttcgcc atctctccat ctctctttct ctctgagata 420
gtctttcgaa tccatctcta gggctcttgt ttctcccat cctccccca cccaccccc 480
caccaaacc aagtcccctt gttcaatccg acaagacaag catcc 525

<210> 24
<211> 587
<212> DNA
<213> Zea mays

<220>
<221> promoter
<222> (1) ... (587)
<223> ESR promoter

<400> 24
gaattcgccc ttggtagatg tctagatgac ctattctact tttcctaaga ttttctctgt 60
atgagtaacc tgtcataatt taacttgtga gatcttgccg atataaaaaa aaaacgccag 120
tcatttatgg tacgggatta ataggttcca agaaccagcc acaatccatt tattagtttc 180
atataaatgt cataaatttt tactaaaatt ttctctgtat agtaacatgt cataactgaa 240
cttgtgagaa aaacgccagt tatttatggg acgggattaa taggttccaa aaaccagccg 300
taacctatth atattagggg actttaagct ggtgccctca gttttgttgg tgtcttcgtt 360
tttaaaactta gttgtattht ttttcttagt tctgtccttc tagtggtata gagcataagg 420
acaaaattga gcaaaaaatg actaaggata aaaatgagga tatcagaaag ggcagcagct 480
taaaaaacct tttatattag ttcaaaagga caccagtcta taaaaagtat actccaagca 540
catttgaaatt tggatttgca ttgtcagtcg ggcagtcgaa ggggacc 587

<210> 25
<211> 900
<212> DNA
<213> Zea mays

<220>
<221> promoter
<222> (1) ... (900)
<223> PCNA2 promoter

<400> 25

```
atcgtaatcg gttttcacccg tataccgaac cgaaaaaacc gaataccaaa ctttatcaat 60
tcccaaattt gactattcga ttatgtgaac taattgtgtg atacaattaa attgttattc 120
acttatttgt atgtgatgta tgatgtatat ctaaataattt gtacctatat aattttttact 180
ttttaaaatt atatgtaatc tatcatgtaa acttggttgta tgtattgtct tgattataag 240
tttggtattc gggtttttacc gaaaaatcga agtaaaaaaac cgaaaccgaa cttctcgggt 300
tttcattttc tagaaaaccg aacgggtttct aatgtttgaa aaaccgaagt tttttaaaac 360
cgaaaaaccg aaccgaagtt tagaaaaaaa ccgaatgccc agccctaaaa attagtaccc 420
cataagaact aaaaaaagat aaaatgacta aaaattaatc agttgaaacc aaacctattt 480
tccccacac ctcacggtat tgtttcgcat tccaagtttg aaacacgact ggaaacaaaa 540
cccaaaacga ctggagggac cgagcttgtg ctgagcagca gagatggcgg gaaatgctgc 600
gtctcccgcc tcagtttcgg atgccccgcc ctttcccaaa ccggccaccg ccgccgcccg 660
tgtctcccca ccgacaggtg ggtccaatcc ttaaccacgg accagggccc ccacctgtca 720
ggtggacctt ccgaagcaag gatcggccag gcgggaaaac atttcgcggc aggtggcggt 780
tgcgccaaat ttctccctcc cttttccggt cggcgtcccc aaacgcctcc ctattaatct 840
ccccgcgttc cccttccctc gcgcgcgcgc tctccctcc caaagctcgc cccgctccca 900
```

<210> 26

<211> 1560

<212> DNA

<213> Zea mays

<220>

<221> CDS

<222> (1)... (1560)

<400> 26

```
atg aag ccg cca tca ctg gtg cac tgc ttc aag ctg ctg gtc ctg ctg 48
Met Lys Pro Pro Ser Leu Val His Cys Phe Lys Leu Leu Val Leu Leu
  1             5             10             15

gcg ctc gcc agg ctg acc atg cac gtc ccc gac gag gac atg cta tcg 96
Ala Leu Ala Arg Leu Thr Met His Val Pro Asp Glu Asp Met Leu Ser
          20             25             30

ccc ctc gcc gcg ctg cgc ctc gac ggt cat ttc agc ttc cat gac gtc 144
Pro Leu Gly Ala Leu Arg Leu Asp Gly His Phe Ser Phe His Asp Val
          35             40             45

tcc gcc atg gcg cgg gac ttc gcc aac cag tgc agc ttc ctg ccg gcc 192
Ser Ala Met Ala Arg Asp Phe Gly Asn Gln Cys Ser Phe Leu Pro Ala
          50             55             60

gcc gtg ctc cac cca gcc tcg gtc tcc gat atc gcc gcc acc gtg agg 240
Ala Val Leu His Pro Gly Ser Val Ser Asp Ile Ala Ala Thr Val Arg
          65             70             75             80

cac gtc ttc tcc ctg gcc gag gcc tcg ccg ctc acc gtc gcg gcg cgc 288
His Val Phe Ser Leu Gly Glu Gly Ser Pro Leu Thr Val Ala Ala Arg
          85             90             95

ggg cat gga cac tcc ctc atg ggt cag tcc cag gcc gcc cag ggg atc 336
Gly His Gly His Ser Leu Met Gly Gln Ser Gln Ala Ala Gln Gly Ile
          100            105            110

gtg gtc agg atg gag tcg ctc cgg gcc gct agg ctc cag gtc cac gac 384
```

Val	Val	Arg	Met	Glu	Ser	Leu	Arg	Gly	Ala	Arg	Leu	Gln	Val	His	Asp		
		115					120					125					
ggc	ttt	gtc	gat	gcc	ccc	gga	gga	gag	ctc	tgg	atc	aat	gtc	ctg	cgt	432	
Gly	Phe	Val	Asp	Ala	Pro	Gly	Gly	Glu	Leu	Trp	Ile	Asn	Val	Leu	Arg		
	130					135					140						
gag	acg	ctg	aag	cac	ggc	ctg	gca	ccc	aag	tcg	tgg	acg	gac	tat	ctc	480	
Glu	Thr	Leu	Lys	His	Gly	Leu	Ala	Pro	Lys	Ser	Trp	Thr	Asp	Tyr	Leu		
145					150					155					160		
cat	ctc	acg	gtc	ggc	ggc	acc	ttg	tct	aat	gcg	ggg	gtc	agc	ggc	cag	528	
His	Leu	Thr	Val	Gly	Gly	Thr	Leu	Ser	Asn	Ala	Gly	Val	Ser	Gly	Gln		
				165					170					175			
gcg	ttc	cgc	cac	gga	ccg	cag	gtc	agc	aat	gtc	aat	caa	ctg	gag	att	576	
Ala	Phe	Arg	His	Gly	Pro	Gln	Val	Ser	Asn	Val	Asn	Gln	Leu	Glu	Ile		
			180					185					190				
gtg	aca	gga	agg	gga	gac	gtc	ggt	acc	tgc	tca	ccc	gag	gat	aac	tct	624	
Val	Thr	Gly	Arg	Gly	Asp	Val	Val	Thr	Cys	Ser	Pro	Glu	Asp	Asn	Ser		
	195					200						205					
gat	ctc	ttc	tat	gct	gct	ctc	ggc	ggt	ctt	ggt	cag	ttc	ggg	atc	ata	672	
Asp	Leu	Phe	Tyr	Ala	Ala	Leu	Gly	Gly	Leu	Gly	Gln	Phe	Gly	Ile	Ile		
	210					215					220						
acc	aga	gca	agg	att	gca	ctt	gag	cct	gct	cca	gag	atg	gtg	agg	tgg	720	
Thr	Arg	Ala	Arg	Ile	Ala	Leu	Glu	Pro	Ala	Pro	Glu	Met	Val	Arg	Trp		
225					230					235					240		
ata	aga	gtt	ctt	tac	tcg	gat	ttt	gaa	agc	ttc	acc	gaa	gac	cag	gag	768	
Ile	Arg	Val	Leu	Tyr	Ser	Asp	Phe	Glu	Ser	Phe	Thr	Glu	Asp	Gln	Glu		
				245					250					255			
atg	ttg	atc	atg	gca	gag	aac	tcc	ttt	gac	tac	att	gaa	ggc	ttt	gtc	816	
Met	Leu	Ile	Met	Ala	Glu	Asn	Ser	Phe	Asp	Tyr	Ile	Glu	Gly	Phe	Val		
			260					265					270				
atc	ata	aac	agg	aca	ggc	atc	ctc	aac	aac	tgg	agg	gcg	tcc	ttc	aag	864	
Ile	Ile	Asn	Arg	Thr	Gly	Ile	Leu	Asn	Asn	Trp	Arg	Ala	Ser	Phe	Lys		
		275				280						285					
cca	cag	gac	cca	gtc	caa	gca	agc	cat	ttc	cag	tca	gat	gga	aga	gtg	912	
Pro	Gln	Asp	Pro	Val	Gln	Ala	Ser	His	Phe	Gln	Ser	Asp	Gly	Arg	Val		
	290					295					300						
cta	tac	tgc	ctc	gaa	cta	acc	aag	aac	ttc	aat	agt	ggc	gac	act	gat	960	
Leu	Tyr	Cys	Leu	Glu	Leu	Thr	Lys	Asn	Phe	Asn	Ser	Gly	Asp	Thr	Asp		
305					310					315					320		
acc	atg	gaa	cag	gaa	gtt	gct	gta	ctg	cta	tct	cgg	ctt	aga	ttc	ata	1008	
Thr	Met	Glu	Gln	Glu	Val	Ala	Val	Leu	Leu	Ser	Arg	Leu	Arg	Phe	Ile		
				325				330						335			
cag	tct	act	cta	ttc	cac	acc	gat	gtc	acg	tac	ctg	gag	ttt	ttg	gac	1056	
Gln	Ser	Thr	Leu	Phe	His	Thr	Asp	Val	Thr	Tyr	Leu	Glu	Phe	Leu	Asp		
			340					345					350				

agg gtg cac acc tct gag ctg aag ctg agg gca caa agc ctc tgg gaa	1104
Arg Val His Thr Ser Glu Leu Lys Leu Arg Ala Gln Ser Leu Trp Glu	
355 360 365	
ggt cca cac cct tgg ttg aat ctt ctg ata ccg agg agc tca atc cgc	1152
Val Pro His Pro Trp Leu Asn Leu Leu Ile Pro Arg Ser Ser Ile Arg	
370 375 380	
aga ttt gct acg gaa gtc ttt ggc agg atc ctg aaa gat agc aac aat	1200
Arg Phe Ala Thr Glu Val Phe Gly Arg Ile Leu Lys Asp Ser Asn Asn	
385 390 395 400	
ggt cct ata ttg ctt tat cca gtg aac aaa tca aag tgg gac aac aaa	1248
Gly Pro Ile Leu Leu Tyr Pro Val Asn Lys Ser Lys Trp Asp Asn Lys	
405 410 415	
acg tca gtg gtc ata cca gat gag gaa att ttc tac cta gtg gga ttc	1296
Thr Ser Val Val Ile Pro Asp Glu Glu Ile Phe Tyr Leu Val Gly Phe	
420 425 430	
ctt tct tca gca ccg tct ctc tca ggt cac ggc agc att gca cat gcg	1344
Leu Ser Ser Ala Pro Ser Leu Ser Gly His Gly Ser Ile Ala His Ala	
435 440 445	
atg agc ctg aac agc caa ata gta gag ttc tgt gaa gag gct gat att	1392
Met Ser Leu Asn Ser Gln Ile Val Glu Phe Cys Glu Glu Ala Asp Ile	
450 455 460	
ggg atg aaa cag tat cta gca cac tac acc aca cag gag cag tgg aaa	1440
Gly Met Lys Gln Tyr Leu Ala His Tyr Thr Thr Gln Glu Gln Trp Lys	
465 470 475 480	
acc cac ttt gga gca agg tgg gag aca ttt gaa cgg agg aaa cac aga	1488
Thr His Phe Gly Ala Arg Trp Glu Thr Phe Glu Arg Arg Lys His Arg	
485 490 495	
tat gat ccc cta gcc atc cta gca cca gga cag aga ata ttc cca aag	1536
Tyr Asp Pro Leu Ala Ile Leu Ala Pro Gly Gln Arg Ile Phe Pro Lys	
500 505 510	
gcg tca ctc cca ttg tct ttg tga	1560
Ala Ser Leu Pro Leu Ser Leu *	
515	

<210> 27
 <211> 519
 <212> PRT
 <213> Zea mays

<400> 27
 Met Lys Pro Pro Ser Leu Val His Cys Phe Lys Leu Leu Val Leu Leu
 1 5 10 15
 Ala Leu Ala Arg Leu Thr Met His Val Pro Asp Glu Asp Met Leu Ser
 20 25 30
 Pro Leu Gly Ala Leu Arg Leu Asp Gly His Phe Ser Phe His Asp Val
 35 40 45

Ala Ser Leu Pro Leu Ser Leu
515

<210> 28
<211> 1617
<212> DNA
<213> Zea mays

<220>
<221> CDS
<222> (1)...(1617)

```

<400> 28
atg gca aga agg act cgt ttc gtg gcc atc gcc gcc ctc ctc aca agc 48
Met Ala Arg Arg Thr Arg Phe Val Ala Ile Ala Ala Leu Leu Thr Ser
 1          5          10          15

ttc ctc aac gtc gca gcc ggg cat tcc cgg cca ctg tcc ggt gcc ggc 96
Phe Leu Asn Val Ala Ala Gly His Ser Arg Pro Leu Ser Gly Ala Gly
      20          25          30

ctc ccg ggc gat ctt ttc ggg ctg ggc atc gcg tcg agg atc cgc acg 144
Leu Pro Gly Asp Leu Phe Gly Leu Gly Ile Ala Ser Arg Ile Arg Thr
      35          40          45

gac agc aac tcg acg gcg aag gcg gcg acg gac ttc ggc cag atg gtg 192
Asp Ser Asn Ser Thr Ala Lys Ala Ala Thr Asp Phe Gly Gln Met Val
      50          55          60

agg gcc gcg ccg gag gcc gtg ttc cac ccc gcc acg ccg gcc gac atc 240
Arg Ala Ala Pro Glu Ala Val Phe His Pro Ala Thr Pro Ala Asp Ile
      65          70          75          80

gcc gcg ctc gtc cgg ttc tcc gcc acg tcg gcg gcg ccg ttc ccc gtt 288
Ala Ala Leu Val Arg Phe Ser Ala Thr Ser Ala Ala Pro Phe Pro Val
      85          90          95

gcg ccg cgc ggg cag gcc cac tcc tgg cgc gcc cag gcg ctc gcc ccg 336
Ala Pro Arg Gly Gln Gly His Ser Trp Arg Gly Gln Ala Leu Ala Pro
      100          105          110

ggc gcc gtc gtc gtg gac atg gcc tcg ctg ggg cgc gcc ccc cgc atc 384
Gly Gly Val Val Val Asp Met Gly Ser Leu Gly Arg Gly Pro Arg Ile
      115          120          125

aac gtg tcc gcc gtg gcc gcc gcg gag ccg ttc gtc gac gcc gcc ggc 432
Asn Val Ser Ala Val Ala Gly Ala Glu Pro Phe Val Asp Ala Gly Gly
      130          135          140

gag cag ctg tgg gtc gac gtc ctc cgc gcc acg ctg cga cac gcc ctg 480
Glu Gln Leu Trp Val Asp Val Leu Arg Ala Thr Leu Arg His Gly Leu
      145          150          155          160

gcg ccc cgc gtg tgg acc gac tac ctc cgg ctc acc gtc gcc gcc acc 528
Ala Pro Arg Val Trp Thr Asp Tyr Leu Arg Leu Thr Val Gly Gly Thr
      165          170          175

```

ctc tcc aac gcg gga atc ggc ggg cag gcg ttc cga cac ggt ccg cag	576
Leu Ser Asn Ala Gly Ile Gly Gly Gln Ala Phe Arg His Gly Pro Gln	
180 185 190	
atc gcc aac gtg cat gaa ctc gac gtc gtc aca ggc aca ggt gag atg	624
Ile Ala Asn Val His Glu Leu Asp Val Val Thr Gly Thr Gly Glu Met	
195 200 205	
gtg aca tgc tcc atg gac gtg aac tcg gac ctg ttc atg gcg gct cta	672
Val Thr Cys Ser Met Asp Val Asn Ser Asp Leu Phe Met Ala Ala Leu	
210 215 220	
ggc ggg tta ggc cag ttc ggg gtc ata acc aga gca cgg atc cgg ctt	720
Gly Gly Leu Gly Gln Phe Gly Val Ile Thr Arg Ala Arg Ile Arg Leu	
225 230 235 240	
gag ccg gcg ccc aag agg gtg cgc tgg gtt cga ctt gcc tac acc gac	768
Glu Pro Ala Pro Lys Arg Val Arg Trp Val Arg Leu Ala Tyr Thr Asp	
245 250 255	
gtc gct act ttc acc aag gat cag gag ttt ctc ata tca aac cgg gct	816
Val Ala Thr Phe Thr Lys Asp Gln Glu Phe Leu Ile Ser Asn Arg Ala	
260 265 270	
agc caa gtc ggg ttc gac tac gtc gaa ggc cag gtc cag ctc agc cgg	864
Ser Gln Val Gly Phe Asp Tyr Val Glu Gly Gln Val Gln Leu Ser Arg	
275 280 285	
tcc ttg gtc gaa ggc ccc aaa tca aca ccc ttc ttc tcc ggc gcc gat	912
Ser Leu Val Glu Gly Pro Lys Ser Thr Pro Phe Phe Ser Gly Ala Asp	
290 295 300	
gtt gct agg ctt gct gga ctc gcg tcc agg acc gga cct gct gca atc	960
Val Ala Arg Leu Ala Gly Leu Ala Ser Arg Thr Gly Pro Ala Ala Ile	
305 310 315 320	
tac tac atc gaa ggc gcc atg tac tac acc aag gac acc gcc ata tct	1008
Tyr Tyr Ile Glu Gly Ala Met Tyr Tyr Thr Lys Asp Thr Ala Ile Ser	
325 330 335	
gtg gac aag aaa atg aag gca ctc ctg gat cag ctg agc ttc gag cca	1056
Val Asp Lys Lys Met Lys Ala Leu Leu Asp Gln Leu Ser Phe Glu Pro	
340 345 350	
ggg ttt gcg ttc acc aag gac gtg acg ttc gtg cag ttc ctc gat cgg	1104
Gly Phe Ala Phe Thr Lys Asp Val Thr Phe Val Gln Phe Leu Asp Arg	
355 360 365	
gtg cgc gag gag gag agg gtg ctc cgg tca gcc ggc gcg tgg gag gtg	1152
Val Arg Glu Glu Glu Arg Val Leu Arg Ser Ala Gly Ala Trp Glu Val	
370 375 380	
ccg cac cca tgg ctg aac ctc ttc gtc cca cgg tcg cgc atc ctc gac	1200
Pro His Pro Trp Leu Asn Leu Phe Val Pro Arg Ser Arg Ile Leu Asp	
385 390 395 400	
ttc gac gac gga gtg ttc aag gct ctg ctc aag gac tcc aac cca gct	1248
Phe Asp Asp Gly Val Phe Lys Ala Leu Leu Lys Asp Ser Asn Pro Ala	

405						410						415						
ggg atc atc ctc atg tac ccc atg aac aag gat agg tgg gac gac cgg	1296																	
Gly Ile Ile Leu Met Tyr Pro Met Asn Lys Asp Arg Trp Asp Asp Arg																		
420 425 430																		
atg aca gcg atg acc cca gcc acg gac gac gac gac atg ttc tat gcc	1344																	
Met Thr Ala Met Thr Pro Ala Thr Asp Asp Asp Asp Met Phe Tyr Ala																		
435 440 445																		
gtt agt ttc ctt tgg tca gca ctg tcc gca gac gac gtg ccc cag ctc	1392																	
Val Ser Phe Leu Trp Ser Ala Leu Ser Ala Asp Asp Val Pro Gln Leu																		
450 455 460																		
gag aga tgg aac aag gca gtg ctg gac ttc tgt gat cgg tca gga ata	1440																	
Glu Arg Trp Asn Lys Ala Val Leu Asp Phe Cys Asp Arg Ser Gly Ile																		
465 470 475 480																		
gaa tgc aag cag tac ctg cca cac tac aca tct caa gac ggg tgg cga	1488																	
Glu Cys Lys Gln Tyr Leu Pro His Tyr Thr Ser Gln Asp Gly Trp Arg																		
485 490 495																		
cgg cat ttc ggg gcg aaa tgg agc agg atc gct gag ctg aag gcc aga	1536																	
Arg His Phe Gly Ala Lys Trp Ser Arg Ile Ala Glu Leu Lys Ala Arg																		
500 505 510																		
tat gac cct cgg gca ttg ttg tcg ccg ggc cag agg att ttt ccg gtg	1584																	
Tyr Asp Pro Arg Ala Leu Leu Ser Pro Gly Gln Arg Ile Phe Pro Val																		
515 520 525																		
cca gta gag gca tct ggc att gct tct gcc tga	1617																	
Pro Val Glu Ala Ser Gly Ile Ala Ser Ala *																		
530 535																		

<210> 29
 <211> 538
 <212> PRT
 <213> Zea mays

<400> 29

Met Ala Arg Arg Thr Arg Phe Val Ala Ile Ala Ala Leu Leu Thr Ser	
1 5 10 15	
Phe Leu Asn Val Ala Ala Gly His Ser Arg Pro Leu Ser Gly Ala Gly	
20 25 30	
Leu Pro Gly Asp Leu Phe Gly Leu Gly Ile Ala Ser Arg Ile Arg Thr	
35 40 45	
Asp Ser Asn Ser Thr Ala Lys Ala Ala Thr Asp Phe Gly Gln Met Val	
50 55 60	
Arg Ala Ala Pro Glu Ala Val Phe His Pro Ala Thr Pro Ala Asp Ile	
65 70 75 80	
Ala Ala Leu Val Arg Phe Ser Ala Thr Ser Ala Ala Pro Phe Pro Val	
85 90 95	
Ala Pro Arg Gly Gln Gly His Ser Trp Arg Gly Gln Ala Leu Ala Pro	
100 105 110	
Gly Gly Val Val Val Asp Met Gly Ser Leu Gly Arg Gly Pro Arg Ile	
115 120 125	
Asn Val Ser Ala Val Ala Gly Ala Glu Pro Phe Val Asp Ala Gly Gly	

130		135		140
Glu Gln Leu Trp Val Asp Val Leu Arg Ala Thr Leu Arg His Gly Leu				
145		150		155
Ala Pro Arg Val Trp Thr Asp Tyr Leu Arg Leu Thr Val Gly Gly Thr				160
	165		170	175
Leu Ser Asn Ala Gly Ile Gly Gly Gln Ala Phe Arg His Gly Pro Gln				
	180		185	190
Ile Ala Asn Val His Glu Leu Asp Val Val Thr Gly Thr Gly Glu Met				
	195	200		205
Val Thr Cys Ser Met Asp Val Asn Ser Asp Leu Phe Met Ala Ala Leu				
	210	215		220
Gly Gly Leu Gly Gln Phe Gly Val Ile Thr Arg Ala Arg Ile Arg Leu				
225		230		235
Glu Pro Ala Pro Lys Arg Val Arg Trp Val Arg Leu Ala Tyr Thr Asp				240
	245		250	255
Val Ala Thr Phe Thr Lys Asp Gln Glu Phe Leu Ile Ser Asn Arg Ala				
	260		265	270
Ser Gln Val Gly Phe Asp Tyr Val Glu Gly Gln Val Gln Leu Ser Arg				
	275	280		285
Ser Leu Val Glu Gly Pro Lys Ser Thr Pro Phe Phe Ser Gly Ala Asp				
	290	295		300
Val Ala Arg Leu Ala Gly Leu Ala Ser Arg Thr Gly Pro Ala Ala Ile				
305		310		315
Tyr Tyr Ile Glu Gly Ala Met Tyr Tyr Thr Lys Asp Thr Ala Ile Ser				
	325		330	335
Val Asp Lys Lys Met Lys Ala Leu Leu Asp Gln Leu Ser Phe Glu Pro				
	340		345	350
Gly Phe Ala Phe Thr Lys Asp Val Thr Phe Val Gln Phe Leu Asp Arg				
	355		360	365
Val Arg Glu Glu Glu Arg Val Leu Arg Ser Ala Gly Ala Trp Glu Val				
	370	375		380
Pro His Pro Trp Leu Asn Leu Phe Val Pro Arg Ser Arg Ile Leu Asp				
385		390		395
Phe Asp Asp Gly Val Phe Lys Ala Leu Leu Lys Asp Ser Asn Pro Ala				
	405		410	415
Gly Ile Ile Leu Met Tyr Pro Met Asn Lys Asp Arg Trp Asp Asp Arg				
	420		425	430
Met Thr Ala Met Thr Pro Ala Thr Asp Asp Asp Asp Met Phe Tyr Ala				
	435		440	445
Val Ser Phe Leu Trp Ser Ala Leu Ser Ala Asp Asp Val Pro Gln Leu				
	450		455	460
Glu Arg Trp Asn Lys Ala Val Leu Asp Phe Cys Asp Arg Ser Gly Ile				
465		470		475
Glu Cys Lys Gln Tyr Leu Pro His Tyr Thr Ser Gln Asp Gly Trp Arg				
	485		490	495
Arg His Phe Gly Ala Lys Trp Ser Arg Ile Ala Glu Leu Lys Ala Arg				
	500		505	510
Tyr Asp Pro Arg Ala Leu Leu Ser Pro Gly Gln Arg Ile Phe Pro Val				
	515		520	525
Pro Val Glu Ala Ser Gly Ile Ala Ser Ala				
	530		535	

<210> 30
 <211> 1566
 <212> DNA
 <213> Zea mays

<220>
 <221> CDS
 <222> (1)...(1566)

<400> 30

atg atg ctc gcg tac atg gac cgc gcg acg gcg gcc gcc gag cca gag	48
Met Met Leu Ala Tyr Met Asp Arg Ala Thr Ala Ala Ala Glu Pro Glu	
1 5 10 15	
gac gcc ggc cgc gag ccc gcc acc atg gcg ggc ggg tgc gcg gcg gcg	96
Asp Ala Gly Arg Glu Pro Ala Thr Met Ala Gly Gly Cys Ala Ala Ala	
20 25 30	
gcg acg gat ttc ggc ggg ctg ggg agc gcc atg ccc gcg gcc gtg gtc	144
Ala Thr Asp Phe Gly Gly Leu Gly Ser Ala Met Pro Ala Ala Val Val	
35 40 45	
cgc ccg gcg agc gcg gac gac gtg gcc agc gcc atc cgc gcg gcg gcg	192
Arg Pro Ala Ser Ala Asp Asp Val Ala Ser Ala Ile Arg Ala Ala Ala	
50 55 60	
ctg acg ccg cac ctc acc gtg gcc gcc cgc ggg aac ggg cac tcg gtg	240
Leu Thr Pro His Leu Thr Val Ala Ala Arg Gly Asn Gly His Ser Val	
65 70 75 80	
gcc gcc cag gcc atg gcc gag ggc ggg ctg gtc ctc gac atg cgc tcg	288
Ala Gly Gln Ala Met Ala Glu Gly Gly Leu Val Leu Asp Met Arg Ser	
85 90 95	
ctc gcg gcg ccg tcc cgg cgc gcg cag atg cag ctc gtc gtg cag tgc	336
Leu Ala Ala Pro Ser Arg Arg Ala Gln Met Gln Leu Val Val Gln Cys	
100 105 110	
ccc gac ggc ggc ggc ggc cgc cgc tgc ttc gcc gac gtc ccc ggc ggc	384
Pro Asp Gly Gly Gly Gly Arg Arg Cys Phe Ala Asp Val Pro Gly Gly	
115 120 125	
gcg ctc tgg gag gag gtg ctc cac tgg gcc gtc gac aac cac ggg ctc	432
Ala Leu Trp Glu Glu Val Leu His Trp Ala Val Asp Asn His Gly Leu	
130 135 140	
gcc ccg gcg tcc tgg acg gac tac ctc cgc ctc acc gtg ggc ggc acg	480
Ala Pro Ala Ser Trp Thr Asp Tyr Leu Arg Leu Thr Val Gly Gly Thr	
145 150 155 160	
ctc tcc aat ggc ggc gtc agc ggc cag tcc ttc cgc tac ggg ccc cag	528
Leu Ser Asn Gly Gly Val Ser Gly Gln Ser Phe Arg Tyr Gly Pro Gln	
165 170 175	
gtg tcc aac gtg gcc gag ctc gag gtg gtc acc ggc gac ggc gag cgc	576
Val Ser Asn Val Ala Glu Leu Glu Val Val Thr Gly Asp Gly Glu Arg	
180 185 190	
cgc gtc tgc tcg ccc tcc tcc cac ccg gac ctc ttc ttc gcc gtg ctc	624
Arg Val Cys Ser Pro Ser Ser His Pro Asp Leu Phe Phe Ala Val Leu	
195 200 205	
ggc ggg ctc ggc cag ttt ggc gtc atc acg cgc gcc cgc atc ccg ctc	672

Gly	Gly	Leu	Gly	Gln	Phe	Gly	Val	Ile	Thr	Arg	Ala	Arg	Ile	Pro	Leu		
210						215					220						
cac	agg	gcg	ccc	aag	gcg	gtg	cgg	tgg	acg	cgc	gtg	gtg	tac	gcg	agc	720	
His	Arg	Ala	Pro	Lys	Ala	Val	Arg	Trp	Thr	Arg	Val	Val	Tyr	Ala	Ser		
225					230					235					240		
atc	gcg	gac	tac	acg	gcg	gac	gcg	gag	tgg	ctg	gtg	acg	cgg	ccc	ccc	768	
Ile	Ala	Asp	Tyr	Thr	Ala	Asp	Ala	Glu	Trp	Leu	Val	Thr	Arg	Pro	Pro		
				245					250					255			
gac	gcg	gcg	ttc	gac	tac	gtg	gag	ggc	ttc	gcg	ttc	gtg	aac	agc	gac	816	
Asp	Ala	Ala	Phe	Asp	Tyr	Val	Glu	Gly	Phe	Ala	Phe	Val	Asn	Ser	Asp		
			260					265					270				
gac	ccc	gtg	aac	ggc	tgg	ccg	tcc	gtg	ccc	atc	ccc	ggc	ggc	gcc	cgc	864	
Asp	Pro	Val	Asn	Gly	Trp	Pro	Ser	Val	Pro	Ile	Pro	Gly	Gly	Ala	Arg		
		275				280						285					
ttc	gac	ccg	tcc	ctc	ctc	ccc	gcc	ggc	gcc	ggc	ccc	gtc	ctc	tac	tgc	912	
Phe	Asp	Pro	Ser	Leu	Leu	Pro	Ala	Gly	Ala	Gly	Pro	Val	Leu	Tyr	Cys		
	290					295					300						
ctg	gag	gtg	gcc	ctg	tac	cag	tac	gcg	cac	cgg	ccc	gac	gac	gac	gac	960	
Leu	Glu	Val	Ala	Leu	Tyr	Gln	Tyr	Ala	His	Arg	Pro	Asp	Asp	Asp	Asp		
305					310				315					320			
gag	gag	gac	cag	gcg	gcg	gtg	acc	gtg	agc	cgg	atg	atg	gcg	ccg	ctc	1008	
Glu	Glu	Asp	Gln	Ala	Ala	Val	Thr	Val	Ser	Arg	Met	Met	Ala	Pro	Leu		
			325						330					335			
aag	cac	gtg	cgg	ggc	ctg	gag	ttc	gcg	gcg	gac	gtc	ggg	tac	gtg	gac	1056	
Lys	His	Val	Arg	Gly	Leu	Glu	Phe	Ala	Ala	Asp	Val	Gly	Tyr	Val	Asp		
		340						345					350				
ttc	ctg	tcc	cgc	gtg	aac	cgg	gtg	gag	gag	gag	gcc	cgg	cgc	aac	ggc	1104	
Phe	Leu	Ser	Arg	Val	Asn	Arg	Val	Glu	Glu	Glu	Ala	Arg	Arg	Asn	Gly		
		355				360						365					
agc	tgg	gac	gcg	ccg	cac	ccg	tgg	ctc	aac	ctc	ttc	gtc	tcc	gcg	cgc	1152	
Ser	Trp	Asp	Ala	Pro	His	Pro	Trp	Leu	Asn	Leu	Phe	Val	Ser	Ala	Arg		
	370					375					380						
gac	atc	gcc	gac	ttc	gac	cgc	gcc	gtc	atc	aag	ggc	atg	ctc	gcc	gac	1200	
Asp	Ile	Ala	Asp	Phe	Asp	Arg	Ala	Val	Ile	Lys	Gly	Met	Leu	Ala	Asp		
385					390					395				400			
ggc	atc	gac	ggg	ccc	atg	ctc	gtc	tac	cct	atg	ctc	aag	agc	aag	tgg	1248	
Gly	Ile	Asp	Gly	Pro	Met	Leu	Val	Tyr	Pro	Met	Leu	Lys	Ser	Lys	Trp		
				405					410				415				
gac	ccc	aac	acg	tcg	gtg	gcg	ctg	ccg	gag	ggc	gag	gtc	ttc	tac	ctg	1296	
Asp	Pro	Asn	Thr	Ser	Val	Ala	Leu	Pro	Glu	Gly	Glu	Val	Phe	Tyr	Leu		
			420					425					430				
gtg	gcg	ctg	ctg	cgg	ttc	tgc	cgg	agc	ggc	ggg	ccg	gcg	gtg	gac	gag	1344	
Val	Ala	Leu	Leu	Arg	Phe	Cys	Arg	Ser	Gly	Gly	Pro	Ala	Val	Asp	Glu		
		435					440					445					

ctg	gtg	gcg	cag	aac	ggc	gcc	atc	ctc	cgc	gcc	tgc	cgc	gcc	aac	ggc	1392
Leu	Val	Ala	Gln	Asn	Gly	Ala	Ile	Leu	Arg	Ala	Cys	Arg	Ala	Asn	Gly	
	450					455					460					
tac	gac	tac	aag	gcc	tac	ttc	ccg	agc	tac	cgc	ggc	gag	gcc	gac	tgg	1440
Tyr	Asp	Tyr	Lys	Ala	Tyr	Phe	Pro	Ser	Tyr	Arg	Gly	Glu	Ala	Asp	Trp	
	465				470					475					480	
gcg	cgc	cac	ttc	ggc	gcc	gcc	agg	tgg	agg	cgc	ttc	gtg	gac	cgc	aag	1488
Ala	Arg	His	Phe	Gly	Ala	Ala	Arg	Trp	Arg	Arg	Phe	Val	Asp	Arg	Lys	
				485					490						495	
gcc	cgg	tac	gac	ccg	ctg	gcg	atc	ctc	gcg	ccg	ggc	cag	aag	atc	ttc	1536
Ala	Arg	Tyr	Asp	Pro	Leu	Ala	Ile	Leu	Ala	Pro	Gly	Gln	Lys	Ile	Phe	
			500					505					510			
cct	cgg	gtc	ccg	gcg	tcc	gtc	gcc	gtg	tag							1566
Pro	Arg	Val	Pro	Ala	Ser	Val	Ala	Val	*							
		515					520									

<210> 31
 <211> 521
 <212> PRT
 <213> Zea mays

<400> 31

Met	Met	Leu	Ala	Tyr	Met	Asp	Arg	Ala	Thr	Ala	Ala	Ala	Glu	Pro	Glu	
1				5					10					15		
Asp	Ala	Gly	Arg	Glu	Pro	Ala	Thr	Met	Ala	Gly	Gly	Cys	Ala	Ala	Ala	
		20						25					30			
Ala	Thr	Asp	Phe	Gly	Gly	Leu	Gly	Ser	Ala	Met	Pro	Ala	Ala	Val	Val	
		35				40						45				
Arg	Pro	Ala	Ser	Ala	Asp	Asp	Val	Ala	Ser	Ala	Ile	Arg	Ala	Ala	Ala	
	50				55						60					
Leu	Thr	Pro	His	Leu	Thr	Val	Ala	Ala	Arg	Gly	Asn	Gly	His	Ser	Val	
65				70					75					80		
Ala	Gly	Gln	Ala	Met	Ala	Glu	Gly	Gly	Leu	Val	Leu	Asp	Met	Arg	Ser	
			85					90						95		
Leu	Ala	Ala	Pro	Ser	Arg	Arg	Ala	Gln	Met	Gln	Leu	Val	Val	Gln	Cys	
			100				105						110			
Pro	Asp	Gly	Gly	Gly	Gly	Arg	Arg	Cys	Phe	Ala	Asp	Val	Pro	Gly	Gly	
	115					120						125				
Ala	Leu	Trp	Glu	Glu	Val	Leu	His	Trp	Ala	Val	Asp	Asn	His	Gly	Leu	
	130					135					140					
Ala	Pro	Ala	Ser	Trp	Thr	Asp	Tyr	Leu	Arg	Leu	Thr	Val	Gly	Gly	Thr	
145				150					155						160	
Leu	Ser	Asn	Gly	Gly	Val	Ser	Gly	Gln	Ser	Phe	Arg	Tyr	Gly	Pro	Gln	
			165					170						175		
Val	Ser	Asn	Val	Ala	Glu	Leu	Glu	Val	Val	Thr	Gly	Asp	Gly	Glu	Arg	
		180						185					190			
Arg	Val	Cys	Ser	Pro	Ser	Ser	His	Pro	Asp	Leu	Phe	Phe	Ala	Val	Leu	
		195					200						205			
Gly	Gly	Leu	Gly	Gln	Phe	Gly	Val	Ile	Thr	Arg	Ala	Arg	Ile	Pro	Leu	
	210					215					220					
His	Arg	Ala	Pro	Lys	Ala	Val	Arg	Trp	Thr	Arg	Val	Val	Tyr	Ala	Ser	
225					230					235					240	

Ile	Ala	Asp	Tyr	Thr	Ala	Asp	Ala	Glu	Trp	Leu	Val	Thr	Arg	Pro	Pro	
				245					250					255		
Asp	Ala	Ala	Phe	Asp	Tyr	Val	Glu	Gly	Phe	Ala	Phe	Val	Asn	Ser	Asp	
			260					265					270			
Asp	Pro	Val	Asn	Gly	Trp	Pro	Ser	Val	Pro	Ile	Pro	Gly	Gly	Ala	Arg	
		275					280					285				
Phe	Asp	Pro	Ser	Leu	Leu	Pro	Ala	Gly	Ala	Gly	Pro	Val	Leu	Tyr	Cys	
	290					295					300					
Leu	Glu	Val	Ala	Leu	Tyr	Gln	Tyr	Ala	His	Arg	Pro	Asp	Asp	Asp	Asp	
305					310					315					320	
Glu	Glu	Asp	Gln	Ala	Ala	Val	Thr	Val	Ser	Arg	Met	Met	Ala	Pro	Leu	
			325						330					335		
Lys	His	Val	Arg	Gly	Leu	Glu	Phe	Ala	Ala	Asp	Val	Gly	Tyr	Val	Asp	
		340						345					350			
Phe	Leu	Ser	Arg	Val	Asn	Arg	Val	Glu	Glu	Glu	Ala	Arg	Arg	Asn	Gly	
		355					360					365				
Ser	Trp	Asp	Ala	Pro	His	Pro	Trp	Leu	Asn	Leu	Phe	Val	Ser	Ala	Arg	
	370					375						380				
Asp	Ile	Ala	Asp	Phe	Asp	Arg	Ala	Val	Ile	Lys	Gly	Met	Leu	Ala	Asp	
385					390					395					400	
Gly	Ile	Asp	Gly	Pro	Met	Leu	Val	Tyr	Pro	Met	Leu	Lys	Ser	Lys	Trp	
			405						410					415		
Asp	Pro	Asn	Thr	Ser	Val	Ala	Leu	Pro	Glu	Gly	Glu	Val	Phe	Tyr	Leu	
			420						425				430			
Val	Ala	Leu	Leu	Arg	Phe	Cys	Arg	Ser	Gly	Gly	Pro	Ala	Val	Asp	Glu	
		435					440					445				
Leu	Val	Ala	Gln	Asn	Gly	Ala	Ile	Leu	Arg	Ala	Cys	Arg	Ala	Asn	Gly	
	450					455					460					
Tyr	Asp	Tyr	Lys	Ala	Tyr	Phe	Pro	Ser	Tyr	Arg	Gly	Glu	Ala	Asp	Trp	
465					470					475					480	
Ala	Arg	His	Phe	Gly	Ala	Ala	Arg	Trp	Arg	Arg	Phe	Val	Asp	Arg	Lys	
			485						490					495		
Ala	Arg	Tyr	Asp	Pro	Leu	Ala	Ile	Leu	Ala	Pro	Gly	Gln	Lys	Ile	Phe	
			500					505					510			
Pro	Arg	Val	Pro	Ala	Ser	Val	Ala	Val								
		515					520									

<210> 32

<211> 1629

<212> DNA

<213> Zea mays

<220>

<221> CDS

<222> (1)...(1629)

<400> 32

atg	gag	gtt	gcc	atg	gtc	gtg	agc	gca	aga	gcc	agc	ctg	ctg	atc	ctc	48
Met	Glu	Val	Ala	Met	Val	Val	Ser	Ala	Arg	Ala	Ser	Leu	Leu	Ile	Leu	
1				5				10						15		
gtc	ctc	tcc	ctc	tgc	tct	ccg	tac	aaa	ttc	ata	cag	agc	ccc	atg	gac	96
Val	Leu	Ser	Leu	Cys	Ser	Pro	Tyr	Lys	Phe	Ile	Gln	Ser	Pro	Met	Asp	
			20					25					30			
ctg	ggc	ccc	ctg	aac	ctg	ctc	ccc	acc	acc	agc	acc	gcg	gcc	gcg	tcc	144
Leu	Gly	Pro	Leu	Asn	Leu	Leu	Pro	Thr	Thr	Ser	Thr	Ala	Ala	Ala	Ser	

35					40					45							
agc	gac	ttc	ggc	agg	ata	ctc	ttc	cgc	gcc	ccg	gcc	gcg	gtg	ctg	agg	192	
Ser	Asp	Phe	Gly	Arg	Ile	Leu	Phe	Arg	Ala	Pro	Ala	Ala	Val	Leu	Arg		
50					55					60							
ccc	cag	tcg	ccg	agg	gac	atc	tcc	atg	ctg	ctc	agc	ttc	ctc	tcc	ggc	240	
Pro	Gln	Ser	Pro	Arg	Asp	Ile	Ser	Met	Leu	Leu	Ser	Phe	Leu	Ser	Gly		
65					70					75					80		
tcg	ccc	tcg	ctg	agc	agg	gtc	acg	gtg	gcg	gcc	agg	ggg	gca	ggc	cac	288	
Ser	Pro	Ser	Leu	Ser	Arg	Val	Thr	Val	Ala	Ala	Arg	Gly	Ala	Gly	His		
85					90					95							
tcc	atc	cac	ggg	cag	gcg	cag	gcc	ccg	gac	ggc	att	gtg	gtg	gag	acg	336	
Ser	Ile	His	Gly	Gln	Ala	Gln	Ala	Pro	Asp	Gly	Ile	Val	Val	Glu	Thr		
100					105					110							
cgc	tcc	ttg	ccc	ggc	gag	atg	gag	ttc	cac	cac	gtc	cgc	ggg	gga	ggc	384	
Arg	Ser	Leu	Pro	Gly	Glu	Met	Glu	Phe	His	His	Val	Arg	Gly	Gly	Gly		
115					120					125							
gaa	ggg	cgt	gcc	tcc	tac	gcc	gac	gtg	ggc	ggc	ggg	ggt	ctg	tgg	atc	432	
Glu	Gly	Arg	Ala	Ser	Tyr	Ala	Asp	Val	Gly	Gly	Gly	Val	Leu	Trp	Ile		
130					135					140							
gag	ctc	ctg	gag	cgg	agc	ctg	aag	ctt	ggg	ctg	gct	ccc	agg	tcc	tgg	480	
Glu	Leu	Leu	Glu	Arg	Ser	Leu	Lys	Leu	Gly	Leu	Ala	Pro	Arg	Ser	Trp		
145					150					155					160		
acc	gac	tac	ctc	tac	ctc	act	gtc	ggc	ggg	acg	ctg	tcc	aat	gcc	ggc	528	
Thr	Asp	Tyr	Leu	Tyr	Leu	Thr	Val	Gly	Gly	Thr	Leu	Ser	Asn	Ala	Gly		
165					170					175							
atc	agc	ggg	cag	acg	ttc	aag	cac	ggg	cca	cag	atc	agc	aac	gtc	ctc	576	
Ile	Ser	Gly	Gln	Thr	Phe	Lys	His	Gly	Pro	Gln	Ile	Ser	Asn	Val	Leu		
180					185					190							
cag	ctg	gag	gta	gtc	aca	gga	cga	ggg	gag	att	gtg	gaa	tgc	tca	ccc	624	
Gln	Leu	Glu	Val	Val	Thr	Gly	Arg	Gly	Glu	Ile	Val	Glu	Cys	Ser	Pro		
195					200					205							
agc	aag	gag	gcc	gac	ctg	ttc	aat	gcc	gtc	ctg	gga	ggc	cta	ggc	cag	672	
Ser	Lys	Glu	Ala	Asp	Leu	Phe	Asn	Ala	Val	Leu	Gly	Gly	Leu	Gly	Gln		
210					215					220							
ttc	ggc	atc	ata	acc	agg	gcc	agg	atc	ctg	ctg	cag	gag	gct	ccg	gag	720	
Phe	Gly	Ile	Ile	Thr	Arg	Ala	Arg	Ile	Leu	Leu	Gln	Glu	Ala	Pro	Glu		
225					230					235					240		
aag	gtg	acg	tgg	gtg	agg	gcc	ttc	tac	gac	gac	ttg	ggc	gcc	ttc	acc	768	
Lys	Val	Thr	Trp	Val	Arg	Ala	Phe	Tyr	Asp	Asp	Leu	Gly	Ala	Phe	Thr		
245					250					255							
agg	gac	cag	gag	ctg	ctg	gtg	tcg	att	ccg	gat	tcg	gtg	gac	tac	gtg	816	
Arg	Asp	Gln	Glu	Leu	Leu	Val	Ser	Ile	Pro	Asp	Ser	Val	Asp	Tyr	Val		
260					265					270							

gaa ggg ttc atg gtc ctg aac gag cgg tcc ctc cac agc tcc tcc atc	864
Glu Gly Phe Met Val Leu Asn Glu Arg Ser Leu His Ser Ser Ser Ile	
275 280 285	
gcc ttc ccc gcg agc gtg gac ttc agc ccg gat ttc ggc acc agg agc	912
Ala Phe Pro Ala Ser Val Asp Phe Ser Pro Asp Phe Gly Thr Arg Ser	
290 295 300	
agc cct agg atc tac tac tgc gtc gag ttc gcg gtc cac cac cac cac	960
Ser Pro Arg Ile Tyr Tyr Cys Val Glu Phe Ala Val His His His His	
305 310 315 320	
ggt tac cag cag cag tct cag gcg gcc gtg gag gcc atc tcg agg cgg	1008
Gly Tyr Gln Gln Gln Ser Gln Ala Ala Val Glu Ala Ile Ser Arg Arg	
325 330 335	
atg agc cac atg gcg tcc cag ctg tac agc gtg gag gtg tcc tac ttg	1056
Met Ser His Met Ala Ser Gln Leu Tyr Ser Val Glu Val Ser Tyr Leu	
340 345 350	
gac ttc ctg aac ccg gtc agg atg gag gag gtg agc ctg ccg agc gcc	1104
Asp Phe Leu Asn Arg Val Arg Met Glu Glu Val Ser Leu Arg Ser Ala	
355 360 365	
ggg atg tgg gag gag gtg cac cac ccg tgg ctc aac atg ttc gtg ccc	1152
Gly Met Trp Glu Glu Val His His Pro Trp Leu Asn Met Phe Val Pro	
370 375 380	
aag gcc ggg gtc gct ggc ttc agg gat ctg ctc atg gac aac gtc tcg	1200
Lys Ala Gly Val Ala Gly Phe Arg Asp Leu Leu Met Asp Asn Val Ser	
385 390 395 400	
ccg gat agc ttc cag ggc ctc atc ctc atc tac cca ctc ctc aga gac	1248
Pro Asp Ser Phe Gln Gly Leu Ile Leu Ile Tyr Pro Leu Leu Arg Asp	
405 410 415	
aag tgg gac acc aac acg tcg gtc gtg atc ccg gac tcc ggg ccc acc	1296
Lys Trp Asp Thr Asn Thr Ser Val Val Ile Pro Asp Ser Gly Pro Thr	
420 425 430	
gcg gac gac ccg gtg atg tac gtg gtc ggc atc ctc agg tcc gcg aac	1344
Ala Asp Asp Pro Val Met Tyr Val Val Gly Ile Leu Arg Ser Ala Asn	
435 440 445	
cct ggt cca gaa gaa gac ggt gac ggc tgc tcc cac cgc tgc ctg cac	1392
Pro Gly Pro Glu Glu Asp Gly Asp Gly Cys Ser His Arg Cys Leu His	
450 455 460	
gag ctc ctc cgc agc cac cgc cgg atc gcc gac gcc gcg gag gcg cgc	1440
Glu Leu Leu Arg Ser His Arg Arg Ile Ala Asp Ala Ala Glu Ala Arg	
465 470 475 480	
ctc ggc gcc aag cag tac ctg cct cac cac ccg acc ccg gcc cgc tgg	1488
Leu Gly Ala Lys Gln Tyr Leu Pro His His Pro Thr Pro Ala Arg Trp	
485 490 495	
cag cag cac ctg ggc cgg cgc tgg gag cgc ttc gcg gac cgc aag gcc	1536
Gln Gln His Leu Gly Arg Arg Trp Glu Arg Phe Ala Asp Arg Lys Ala	

500							505					510					
cgg	ttc	gac	ccg	ctg	cgc	atc	ctg	ggg	ccc	ggc	cag	ggc	ata	ttc	cct	1584	
Arg	Phe	Asp	Pro	Leu	Arg	Ile	Leu	Gly	Pro	Gly	Gln	Gly	Ile	Phe	Pro		
515							520					525					
cgg	acg	gcc	cag	gat	gct	gcc	gcc	gct	gct	gcg	tac	ggg	agc	tag	1629		
Arg	Thr	Ala	Gln	Asp	Ala	Ala	Ala	Ala	Ala	Ala	Tyr	Gly	Ser	*			
530							535					540					

<400>	33														
Met	Glu	Val	Ala	Met	Val	Val	Ser	Ala	Arg	Ala	Ser	Leu	Leu	Ile	Leu
1				5					10					15	
Val	Leu	Ser	Leu	Cys	Ser	Pro	Tyr	Lys	Phe	Ile	Gln	Ser	Pro	Met	Asp
			20					25					30		
Leu	Gly	Pro	Leu	Asn	Leu	Leu	Pro	Thr	Thr	Ser	Thr	Ala	Ala	Ala	Ser
		35					40					45			
Ser	Asp	Phe	Gly	Arg	Ile	Leu	Phe	Arg	Ala	Pro	Ala	Ala	Val	Leu	Arg
	50					55				60					
Pro	Gln	Ser	Pro	Arg	Asp	Ile	Ser	Met	Leu	Leu	Ser	Phe	Leu	Ser	Gly
65					70					75					80
Ser	Pro	Ser	Leu	Ser	Arg	Val	Thr	Val	Ala	Ala	Arg	Gly	Ala	Gly	His
				85					90					95	
Ser	Ile	His	Gly	Gln	Ala	Gln	Ala	Pro	Asp	Gly	Ile	Val	Val	Glu	Thr
			100					105					110		
Arg	Ser	Leu	Pro	Gly	Glu	Met	Glu	Phe	His	His	Val	Arg	Gly	Gly	Gly
		115					120					125			
Glu	Gly	Arg	Ala	Ser	Tyr	Ala	Asp	Val	Gly	Gly	Gly	Val	Leu	Trp	Ile
	130					135					140				
Glu	Leu	Leu	Glu	Arg	Ser	Leu	Lys	Leu	Gly	Leu	Ala	Pro	Arg	Ser	Trp
145					150					155					160
Thr	Asp	Tyr	Leu	Tyr	Leu	Thr	Val	Gly	Gly	Thr	Leu	Ser	Asn	Ala	Gly
				165					170					175	
Ile	Ser	Gly	Gln	Thr	Phe	Lys	His	Gly	Pro	Gln	Ile	Ser	Asn	Val	Leu
			180					185					190		
Gln	Leu	Glu	Val	Val	Thr	Gly	Arg	Gly	Glu	Ile	Val	Glu	Cys	Ser	Pro
		195					200					205			
Ser	Lys	Glu	Ala	Asp	Leu	Phe	Asn	Ala	Val	Leu	Gly	Gly	Leu	Gly	Gln
						215					220				
Phe	Gly	Ile	Ile	Thr	Arg	Ala	Arg	Ile	Leu	Leu	Gln	Glu	Ala	Pro	Glu
225					230					235					240
Lys	Val	Thr	Trp	Val	Arg	Ala	Phe	Tyr	Asp	Asp	Leu	Gly	Ala	Phe	Thr
				245					250					255	
Arg	Asp	Gln	Glu	Leu	Leu	Val	Ser	Ile	Pro	Asp	Ser	Val	Asp	Tyr	Val
			260					265					270		
Glu	Gly	Phe	Met	Val	Leu	Asn	Glu	Arg	Ser	Leu	His	Ser	Ser	Ser	Ile
		275					280					285			
Ala	Phe	Pro	Ala	Ser	Val	Asp	Phe	Ser	Pro	Asp	Phe	Gly	Thr	Arg	Ser
		290				295					300				
Ser	Pro	Arg	Ile	Tyr	Tyr	Cys	Val	Glu	Phe	Ala	Val	His	His	His	His
305					310					315					320

aatgcaaaca	aagccatttg	cagaatgtgc	tacatagcag	gtatgtttct	ctttttttcc	1080
ctgtaaaatt	tgtagactta	tcacaagaat	aagtttaacc	attactagaa	tagttcctca	1140
catgtttgtt	taccatcggg	gcgggaacag	cttgcatgtc	aaaagctgcg	caagtattag	1200
ggccctctag	atTTTTTTaa	tagtagtagt	atatataata	tataggtgtt	actatttgag	1260
ttgttaggcc	atctgcggca	gattttctat	gacatccctt	atTTcaaact	ttattttgca	1320
aacagttgtc	atatacccta	TTTTaggcga	atcactgaag	acaggtaagt	tttggcacgg	1380
atgaggtgga	gagtggacaa	gaatctccgt	tgtggagtct	gcctaccagt	accaggcaaa	1440
gtaatgcatg	cgcgcgga	ggatggacgg	tcgaagtggc	ctccctgcct	ccaccccgac	1500
gacgacgcat	gggctccgtc	cccttcgctt	gcttccctgt	ccagctagct	ccatcgccca	1560
gtgctccgct	ccgcccgcaca	ggaacggaac	ggaacggacc	gaaccacttg	gtcgcacccc	1620
gatgcgttgc	cgtctgccgg	tgtccatcgt	gtcgggttca	cctctgcact	agcataaatt	1680
ccttgacacc	aacagcgagc	gacatcatcg	gtcagccctt	acaagtcacg	agtgtttctga	1740
ctgaccagct	agcaatagca	atctgctgct	ctgcttgact	tgtctggacg	atccgcccgt	1800
gcttgcggtt	ggctccagta	ggctatccct	cgcgacgtcg	tcgatctgga	ctccatggcg	1860
tccacacaga	atcgacacga	gcttggtgtg	ccgcgtacgc	atgtgtgctg	atgtatgcct	1920
cgtcttccac	atgcaaacat	acgcagagga	aggggaaagg	cggcagcaaa	cgcgacgggtc	1980
caagtcgtac	cacagaagtg	gtcgcgcatg	tgtgcccaag	ttgccatcac	ccggatgcta	2040
ttagatttcc	agaaactaac	ttgtgaggac	ccctggtgtc	tgctagctgc	tctccaactc	2100
caacctgtca	atcaattccc	agacggacaa	gctgagctca	cagctcaagc	tcaacaacga	2160
tggccggccg	ggtcaccatg	gaactgatcc	tctacagtac	aggcatggga	aaatggagga	2220
ggagagcagg	gcagtgaggc	cacagaatca	gaggctgatt	agtgttggtg	agctccaatc	2280
caacagcata	tgaccagcga	gcagaacata	gggatgtcct	gtgggcttgc	ccagggacag	2340
acgcatgcaa	gccatgtgac	tgtccggaga	gagagccggg	gatactggaa	cagaggatcc	2400
gatectgccc	cccttctttt	gcctctccct	ctctcacaca	cacagtctca	cctatatgtg	2460
gctatgtcgt	ctccattagg	ctgttaacta	gccaacacat	gttccccctg	tgcttaagac	2520
agcagctaca	aagcgagaac	atcatgctct	aaaaagaaac	ttccgcaatg	caccactagc	2580
acatgtctgc	gcctcaattc	gcaaccggca	agcaagcaag	ccggcaagca	gacagtgcgc	2640
atacggtttt	taccaaacag	ctagcgccca	cagctgacta	gctgaccacc	gcaccaccca	2700
cactcctcct	cgcgagtcgc	gaggcaagcc	gcaagctcct	atatagagag	gccccctccc	2760
tccccctgca	tggacagcca	ccgccttctt	caaccctcct	tccgtcttcc	tctcttagtc	2820
ttacctcggt	gcacctcaag	aaacttggcg	cgcaaccagg	aaacccccct	ttctctctct	2880
ctctctctct	ctctctctgc	cttctgattc	caagctcccc	aactgcccag	caccaacctg	2940
ccgaactccc	ctcctttttg	ttggtttgtc	gaattataaa	ttgagccccg	ccggctgact	3000
acc						3003

<210> 35

<211> 2001

<212> DNA

<213> Zea mays

<220>

<221> misc_feature

<222> (0)...(0)

<223> Promoter for ZmCkx3

<221> misc_feature

<222> (0)...(0)

<223> n = A, T, C, or G

<400> 35

ccggggtgtg	acaggagcat	tgaagcatgc	atgctctgct	cagcatataa	ttaaagaaag	60
aagcatcaaa	atgcactgga	gcagttgacc	aaaacttgca	gtacgtcaa	aatatatacg	120
agggctggca	tcaaggtgtg	ctcagcccca	gccccgtcag	gtaacttggt	cttttgTTTT	180
ctggcccttg	ggcttcatta	aaggccgcgc	gccgcgagcg	aggcaaaaca	gtgaagggga	240
ggggaggtgc	cgccactaa	cctctcggtc	ggatatatta	gtattcaagc	agttgacaaa	300
tctgtgctga	tttgatttgg	tctgaggaaa	atatatatat	atatatatat	agccccctgt	360
cgttcatgca	ccctctcgca	gcctgcaacc	ttacaatatt	gttcttgcac	ccggtttttat	420
ttatatTTTT	atTTTTTaaa	aaaaaaatcc	atagtcctgc	cgtcttgaag	gatattgttt	480

tctttaccca	tgcacggcgg	agtttaaatt	tgcgctgacc	cgactgctcg	tgaacagaga	540
caagtatgac	agatatcggt	gagttccaaa	ttttaaaaaa	aaaatcaata	aaaaatttaa	600
aacagaatgt	tgacgaggaa	aaaaaatatg	aagggtgctt	cacacctgtc	actccatgcc	660
ggacatcaac	aaattaattg	ttcaagtggg	gggagtcagc	tgcttccagt	ttaccttcc	720
gcgccagcgg	ttggtagaca	ggattgttgc	cacgtggacg	aaatctcctg	ccgccagctg	780
gttgatcacg	gcaggcagtc	acatgcttct	tgccaagatt	accgcggggt	gtaatcatct	840
gaaatatatt	aacctgagca	cgtgatagag	taaaaaaatt	ggtcgactaa	gggggtgttt	900
ggtttctagg	gactaatgtt	tagtccctac	atttttattcc	attttagttc	taaaattacc	960
aaatatagaa	actaaaactt	tattttagtt	tctatattag	caatttatag	actaaaaaag	1020
aataaaatga	agggactaaa	tattaatccc	tagaaaccaa	acacccccta	acttttaggt	1080
agttgtggca	tgcattctct	ggaacggcag	ttctagagag	cacttgagat	gtcaacaggt	1140
gaagaattga	agattggcca	acacaggcgt	tcaaggagat	tcaaccaccc	atccacatac	1200
cgcgcaaaaca	cttggggggc	attcttgctg	ctgccacatt	tggaagaagc	gcagcaatgt	1260
ggtgttcaga	agaagcacag	ctattttagc	tcttgataac	tatctttttt	tttgcataga	1320
ttaattttatt	tcttcgatat	atactagctt	gtaaaaaaat	gttttncaga	tatatgtata	1380
aaaatgtgta	cctagtacct	acgcatgtct	tagttcaaca	tacttgatag	ctgtagtttt	1440
ctgaaaacct	gttcaaatta	acctttttcc	tacctgatg	gtgaatagag	agaaaagctt	1500
tacctttgtc	tgaataagaa	aactaacaga	aagcttacat	tttggccact	ctacctgccc	1560
gagtattttc	taagcaagca	aaggcgcag	aaaattttct	cggaatccat	gaccttttac	1620
gcgcantgnw	aaayawwgm	mattgmtcmg	accaatgatc	attttgatac	tctccacaag	1680
tcaacatctc	aaaaaaacca	caagatgggg	cccatcaaca	taagttcacg	agtgtgcctt	1740
caggtacatt	gttctttttt	tttgttttgc	taaagtcaat	cagctgcaaa	atattcagaa	1800
caatttcaat	aacccgaaag	gctgttgtgc	ctccatttgc	caacgtttgc	gaggccaaat	1860
ggtacccccg	ctataaatat	catggaagtt	cttggcctct	aggacacaca	agcgatctct	1920
cctcctatag	tttctataac	cccacaaagc	gtccagggtcc	cgtagtcacc	tccgattgca	1980
ttgcgttgcc	gcaagacaag	c				2001

<210> 36

<211> 2448

<212> DNA

<213> Zea mays

<220>

<221> misc_feature

<222> (0)...(0)

<223> promoter for ZmCkx4

<400> 36

ctttatgttg	tagccaagga	aagtatactg	ttaagatcag	aatgaacctt	ataggagtgtg	60
tatgggcata	aagccagcaa	gtatagccaa	aggtacacaa	ggctaataata	gtcaagttgt	120
tgatgtgtga	gacgttcaag	gaagtgaact	attggaggag	tcgactaaaa	gtacgattaa	180
taaggtagac	atgatggtaa	aatctttgat	ctagaattta	agtggtagtg	atgcgagggg	240
gagaatggca	agcacaactt	caaatatagg	gtgatgctta	tgcttggtcg	agccatttca	300
ttcatgagca	taggaacatg	agacatgggtg	ggatatggat	acttgcacaa	aaaaaggaat	360
taagtttatg	atattcacct	cccagtcagt	ttgcatggta	aaaaaattcc	tatcaatttg	420
gttctcaact	agggcctaaa	attctcaaaa	tatctgttgg	ggaccattat	cgtcgacgat	480
cctcagaatc	tgttattacc	aaattaaaaag	gtgtgtttca	ggtactgtgc	aaagcagcag	540
cgaagctatc	cttcgtcaaa	agtggctcaa	tgaaccaggt	ggagaagcta	tgtagcttcg	600
tctgcgtaga	gcgtgccgga	ggaggaagct	ttggctctga	atgcatcgac	ttacgaagca	660
tgggagaaga	agactcagaa	ggcttgtcca	gcgtgggaat	aaaaaggaga	aaatacaatt	720
ttgcccttgt	gggatttgta	aatcatgtgc	aaggctcatg	gatatgtttg	taattttata	780
tgatatgttt	gtaaatcatg	gatatgtttt	gtaaatcagg	tggactagag	gagagggagg	840
gtggacatag	tgacttgcag	cttgatcatg	gtagagtggg	catggtagag	ggaaaggggt	900
aggccaattc	tggagtgcgg	ccacggtgag	ttgagtgtcg	gccacggtag	gggaaagggg	960
tagcccaatt	ctagggccgg	catcgagaaa	ggccgacatg	tgacgctcag	gaggtagtgt	1020
tagagggtttg	aacggaaaaa	attgaacatg	ttagtatgat	gagttgtgta	attgctggga	1080
attgtggata	atttccactt	aactacggcc	ctgtttatatt	acccttagat	tataaaatcc	1140
aacttaaaaa	agttgagatg	taaacaaaca	acacatatta	ttaggtggat	tatgttatct	1200

agaaatctgg	atgataataa	tttataagtc	ggttaatagg	tgtttacata	atcgataagc	1260
tggattatat	aatcctggaa	cacggctttc	gcgagagcgt	attaaaacag	gattccgtga	1320
agcacactat	ctgaggagct	ccaccaaag	ctgaatctag	cccgcactct	tttttggagg	1380
attcaaattt	ggtgtcactg	gagcattcgg	cattttgttt	catggcgtga	agctattttt	1440
actaattaca	gaagctgttt	caaataagacc	tttaaataat	ggctgagtat	aaaaggaggc	1500
aattttttta	tctcgccgat	ggagccaggt	cgcgtcgcgc	cgcggccgtg	ctgcgctctc	1560
gacgcgatct	agcggcgatg	tgcacagtac	agttttgcc	tgccattggt	taagcctgca	1620
tacaacacac	cagcgtactg	ccctgcacaa	gatctcctcg	gctcggcctc	tctgatgga	1680
acgttcagct	tgaacagcgg	agcgtggggg	catcccgggg	atgggcgcgc	cggccgagaa	1740
attttgcaac	ctggcaaata	tgccctgtcg	catactacca	tccacctcca	ggcgccaaga	1800
acgcctccga	gtttcaggct	tgcagctcag	ctctgtgttg	aattggaacg	ggcggagtgt	1860
ctgggttcca	gacttccagt	acaaggcgat	caattggtag	ggcgaattac	ttgcaggccc	1920
agatgcatgg	cccattctatc	tggttctcta	tgcgttgctt	ttacttgac	aatagtggca	1980
gacaaactac	aagtcagatc	cgatcctatc	catccatcca	tctcgcagcg	cgatgcaaat	2040
atgcaatcgt	ctgtggaact	cgaaaaaaa	cagaggctcg	gcctcgcacg	aggttaaggg	2100
aaaaaaaaacg	aagcgttttg	aacttttggt	ggcattcgca	gcatgctgtg	ctgccaccgt	2160
atgtttttat	ttttgctttg	tttgtcttct	ttgagaaacg	tgagggagcc	gcgtgtccgc	2220
tcgttataaa	acccccccgg	cgacccaac	taccacgagc	tcaagcctca	agcctcaagc	2280
ctcaagcaag	cagagcgccg	tgacatcacg	aaacaaacat	atagagctag	ctgctctgcc	2340
tctgcttcac	caatcacctg	cttgcccgcg	cggaggggag	ggtttcccc	tttgacacag	2400
ctgagctccc	ctccatcagc	agccagctcc	tcgtcgcaaa	gcaagaag		2448

<210> 37

<211> 2346

<212> DNA

<213> Zea mays

<220>

<221> misc_feature

<222> (0)...(0)

<223> promoter for ZmCkx5

<400> 37

tacagatttg	cgttcatcaa	tggcagcgcg	ggatctcatg	aggctactgg	gttcttgcaa	60
gtggggagag	aaaggagat	ctacgaaaga	cttgtagtg	ggccaccttt	tccctctttc	120
cccacaagga	cgagatcgtg	gattagagta	ggaaagtgat	tccgcattgg	tctcaaactc	180
tggcgaaaga	ttgcattgtg	tactctccac	cactcgaccg	gcaacgaggg	attttggtat	240
tgcacgatgc	atcctttgca	catgagctag	gcttggtcct	ttgagtattc	agtttagcatt	300
gcaaccccat	ttcaattcac	atgcttgtct	ttccaaggaa	ctttctaagc	cacctaacag	360
acattagggg	ttatatcaga	atcgagctca	tggcgtaact	tatgctgcac	gaacaatggg	420
ttggggggcgt	cgtttcttgc	atgagagcat	gcgcatactg	gtaaggattt	cgccaaaaga	480
acttttagtcc	tctaccgact	ttgtgtttgc	gtgatctcgt	gatttgaagc	ctgtggtggt	540
gtgctgaggg	agcatattgg	aaggatatct	tgtgttgata	tggcatccgt	ccgtggacaa	600
atcgatacca	catactgttc	ttggattcta	ttcttgggat	tgctaaatga	tctagataga	660
ttatattctc	ttgttgacgc	ccctattgct	tcaatacgaa	gaaaacccaa	cgtttagaac	720
ttaataaaaac	catttggtgag	cttagctgct	taggcaattc	atttttatgc	atgacaaata	780
tataataata	ttagctatac	tattattgat	gcaacctgtg	ggagcgtata	aaatgggtact	840
tccccaattc	taaattataa	gacgttttga	ctatatattc	tacatacata	tgtttaattt	900
tatatattaga	taatcgctat	gccttaatat	atagtaaaaa	gtagtatatc	tagaaaagat	960
aaaacatctt	ataatttaaa	aatgggtaga	gtattatatt	agatatgaac	agtgccttaga	1020
tgccaccaa	attttgccat	gccatcctaa	ggccagcaaa	agtttgtgtc	ttcttttgtt	1080
ttccaaacca	ctagatgcc	atatactatt	tatcatcgat	cgagatgtag	gtcttagtta	1140
attgtgtcgg	gtgcccttga	gaaagaaaag	aaaaagggtg	gattttgttt	tcgcttagac	1200
gatgattgga	tctcttgggc	tctgaattcc	atcccgaata	aacaaatgaa	gtaggtcctc	1260
agtcaccctt	gccctgttag	ctgcaagaga	gctcatgggt	tccagccaca	caatcagtc	1320
atggctcctt	cttcttggcc	taagtgggtg	ccaatcttgg	tgggtgatcg	agtcctgggc	1380
cctctgaaca	gtattacaca	acagtaatcc	tgcaaaagat	ttggtatatc	tagattctag	1440
agtgagcgcc	gtgttgtgcc	cagctaggaa	tgggtgttca	agtgcaacag	gaggaggacc	1500

caggatgggc	aggtgtaata	ggctctcatt	aaaagactgt	tcagatggat	tagagcaacg	1560
acggggaagc	cgggaaaaaa	tggttgggtc	tgctttcctc	tcgctccccg	gccgggttca	1620
tatatgaatc	tgagaacgat	atTTTTtTgct	tcatttttca	tttgctatat	atttaaactg	1680
tttttttTgtg	Tgtgtgtgtg	Tgttcattga	gctcaatact	Tgaggcttga	tagggagagg	1740
agtgaggcag	ctgatcacat	ggacctccat	ctgaggacag	ttcctcttcc	gaaacagaaa	1800
ggagagtgcA	gggaccagcg	tggcctgtac	agtattgtgt	ttgccctttt	cctttggcag	1860
ggacagagag	cttcaggctt	gtcctcttta	Tgtatgctgc	tcgcctgctt	cagagtcaga	1920
gcttccccctt	ctcactttctc	agagagagag	agagagaaga	gagagagagg	agagccctcc	1980
acagctcccc	Tgtcctgccc	tcaggcattc	tttgtcacag	ggggcgaggg	ctgaagatca	2040
tcacatgggtg	gcctttttttg	ggTctgtggc	ctttgggtctt	ttagtgtctc	ttcctttttac	2100
ctcctcatga	catgaacccc	ctttttaaac	ctccctcaaa	atcaaatacac	cctccttctc	2160
ctttaagagc	cctcaacccc	ttccctcat	tttccttcat	ccctcagcct	ttgcacaaag	2220
ggcaagaata	acgcagtatg	atcatctgat	catactcccc	ccgccatcac	aatcccacac	2280
gaacgtgaga	caaaggtaac	agacgcaaga	agctagcagc	Tgcaggagat	Tgctcagccc	2340
atctcc						2346

<210> 38

<211> 51

<212> DNA

<213> Zea mays

<400> 38

caucaucauc	auggatccac	caatggatct	acgtctaatt	ttcggtccaa	c	51
------------	------------	------------	------------	------------	---	----

<210> 39

<211> 42

<212> DNA

<213> Zea mays

<400> 39

cuacuacuac	uagttaactc	acattcgaaa	Tggtggctct	tc		42
------------	------------	------------	------------	----	--	----